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FIG. 1

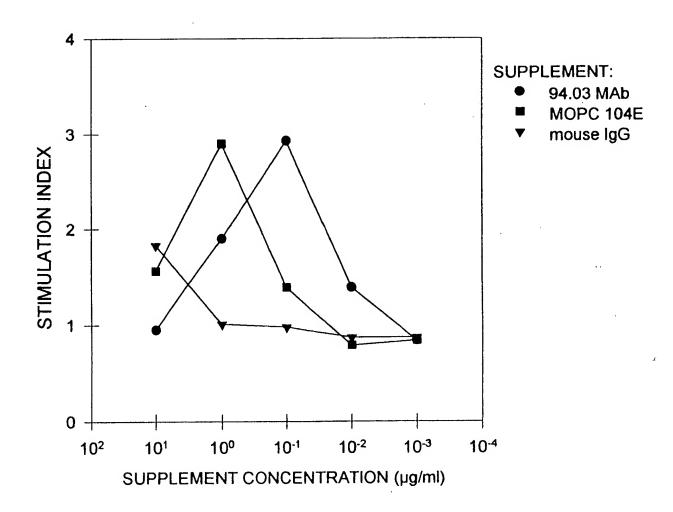
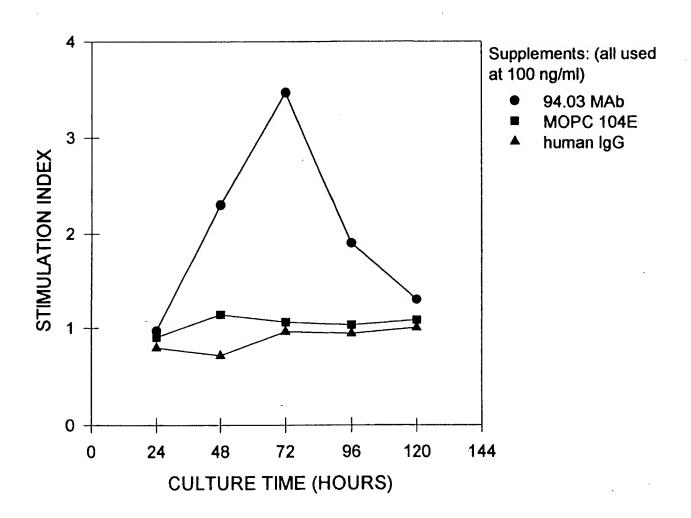




FIG. 2



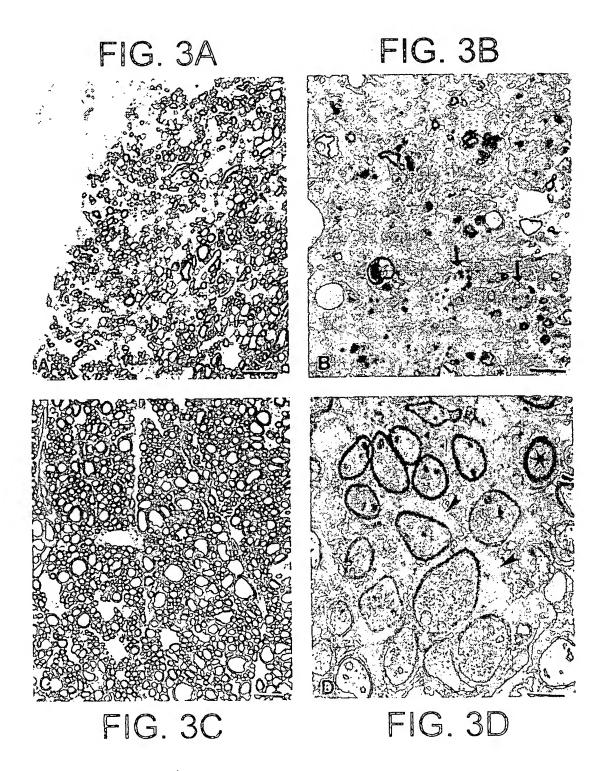
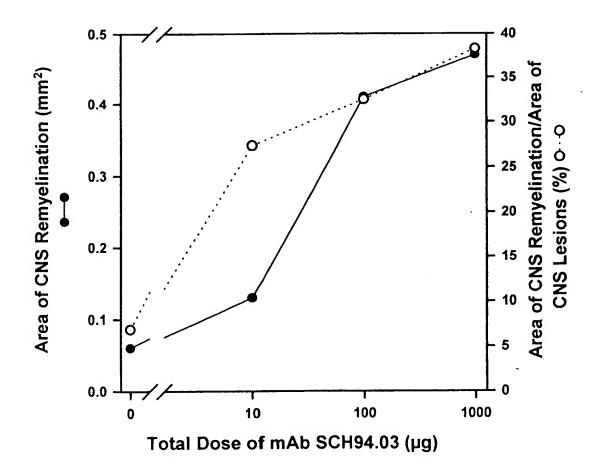
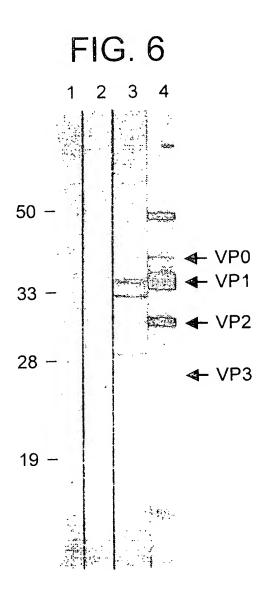


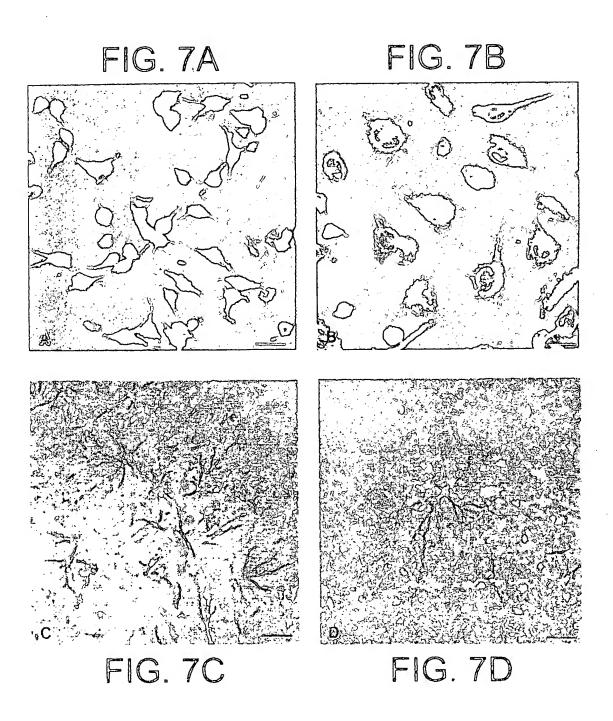


FIG. 5

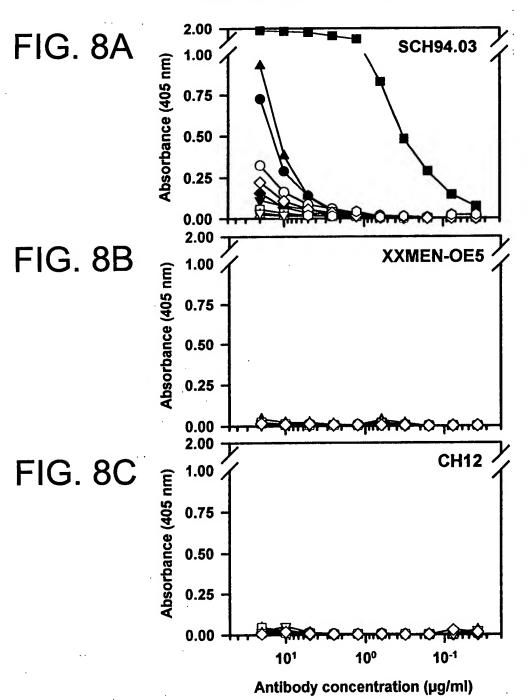








Protein antigen ELISA with SCH94.03



Antigen:

- KLH ■ spectrin
- ▲ hemoglobin
- ▼ vimentin◆ thyroglobulin
- o actin
- □ lysozyme
- △ transferrin
- **▽** myosin
 - tubulin



FIG. 9

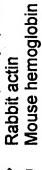
ELISA with SCH94.03 F(ab₂)' fragments

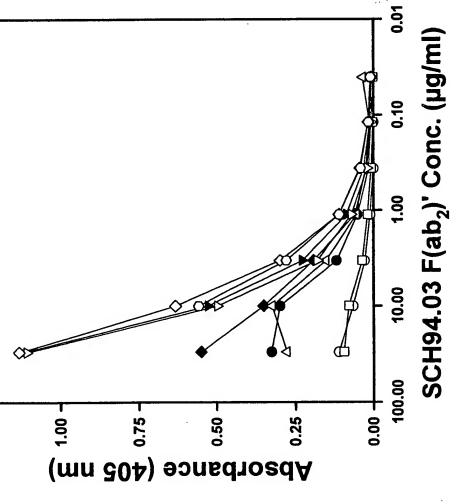
1.25





- Bovine thyroglobulin
 - Human spectrin Bovine tubulin
- Bovine myelin basic protein **Bovine myosin**
 - 0
- **Frinitrophenyl (TNP)-BSA** Dog myoglobin







Chemical hapten ELISA with SCH94.03

FIG. 10A

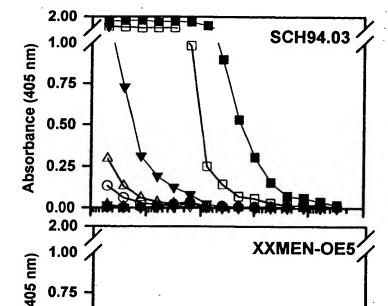
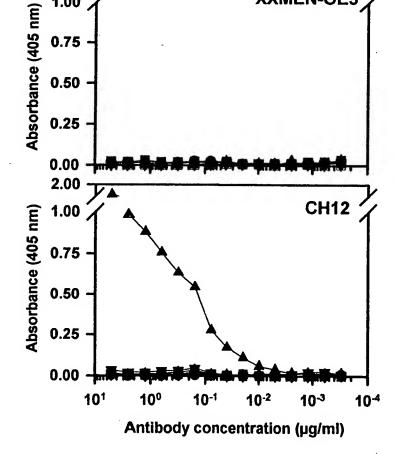


FIG. 10B





Hapten:

- none
- FL
- **▲** TMA **▼** PhOx
- O Ars
- △ TNP▽ PC

16.11A Immunoglobulin Light Chain Variable Region Sequence of SCH94.03

						H	Leader region	H Q	gion															
SCH94.03	M M S ATG ATG TC	s s TCC TCT	A TOOL	o Sa Sas	TIC	H E	o GGT	H CH	r offo	-9 116 0	r DED	C F	T CAA	A GGT	F ACC	C AGA	o t	d d d	AIC	CAG SAG	ATG	H AC H	0 KJ	ACT
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S E S Q : TCA GAG AGT CAG : ---

CAA GGC ACC ACT C

scH94.03 CH12 germline J_H2

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J region

 $J_{\rm H}^2$



Immunoglobulin Heavy Chain Variable Region Sequence of SCH94.03

<u>n</u>						3	Leader region	reg	Lon															
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germline JH3 A2B5



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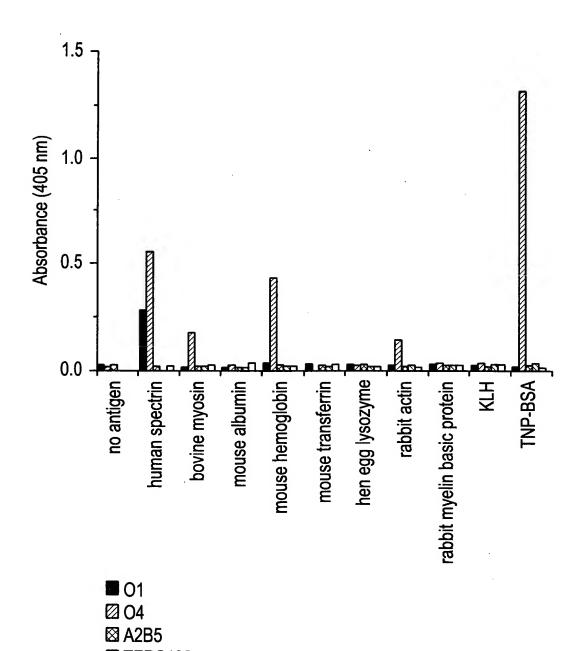


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FIG. 18



☐ TEPC183



FIG. 19A

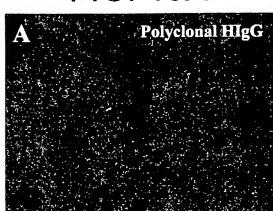


FIG. 19B

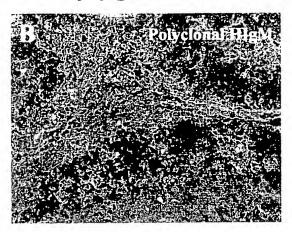


FIG. 19C

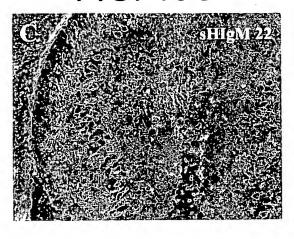


FIG. 19D

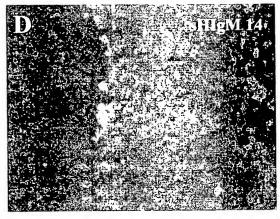


FIG. 19E



FIG. 19F





FIG. 20A

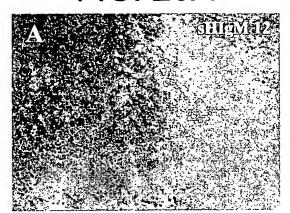


FIG. 20B

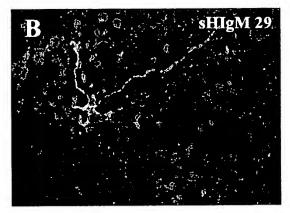


FIG. 20C

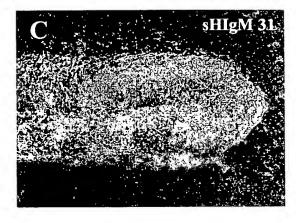


FIG. 20D

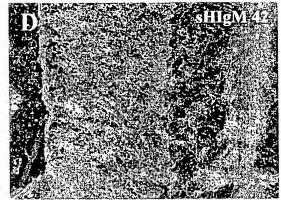


FIG. 20E



FIG. 20F







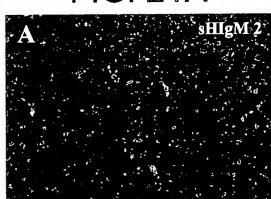


FIG. 21B



FIG. 21C

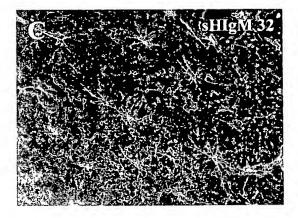


FIG. 21D

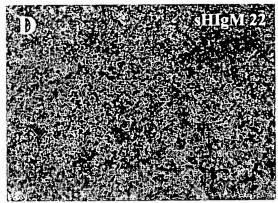
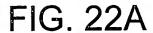


FIG. 21E







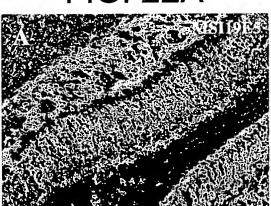


FIG. 22B

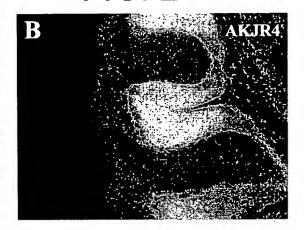


FIG. 22C

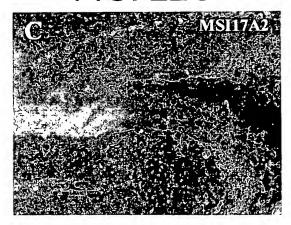


FIG. 22D

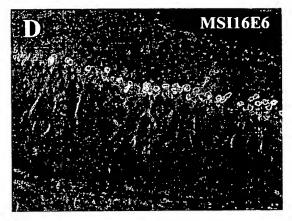


FIG. 22E

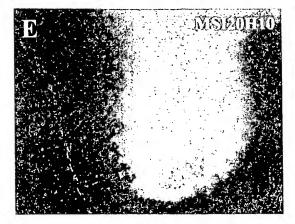


FIG. 22F

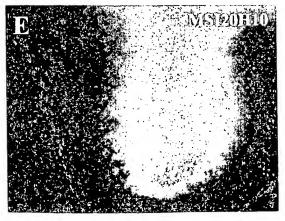








FIG. 23B

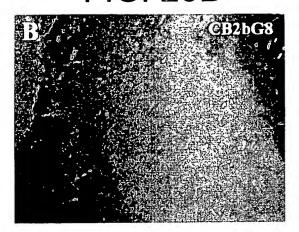


FIG. 23C

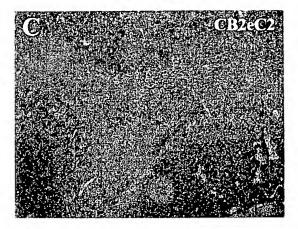


FIG. 23D

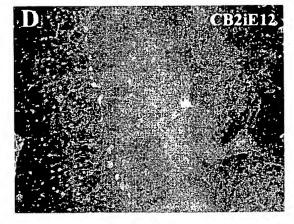


FIG. 23E

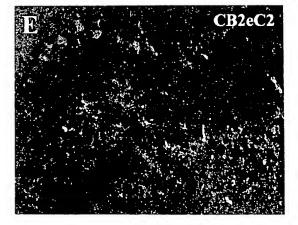
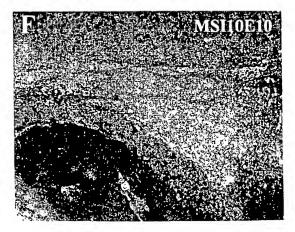
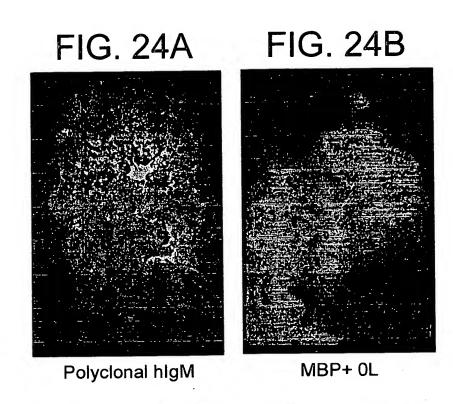


FIG. 23F







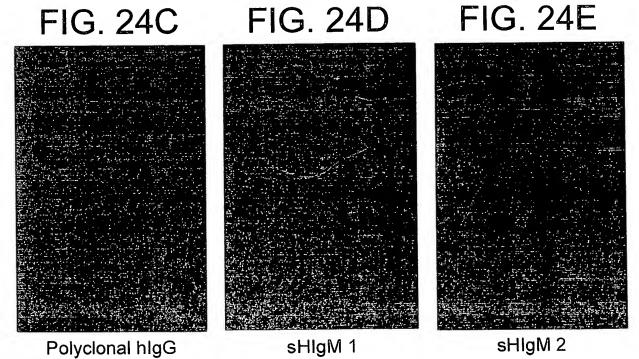






FIG. 25B

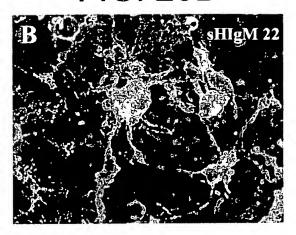


FIG. 25C

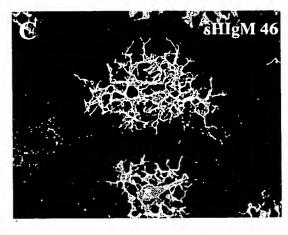


FIG. 25D

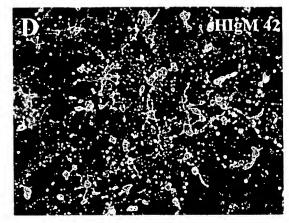


FIG. 25E

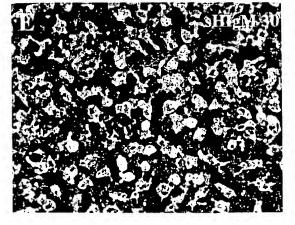


FIG. 25F







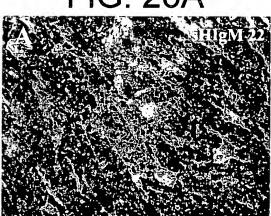


FIG. 26B

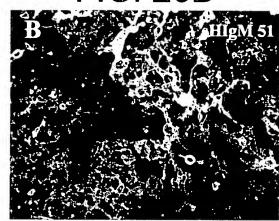


FIG. 26C

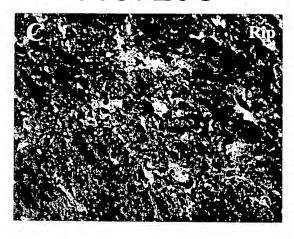


FIG. 26D

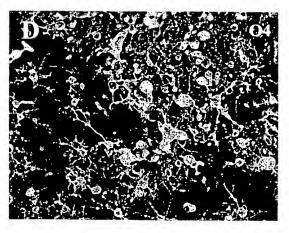


FIG. 26E

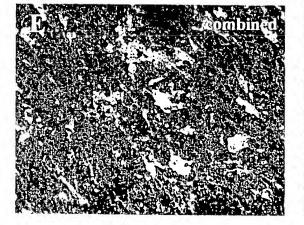
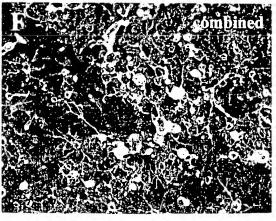
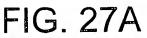


FIG. 26F





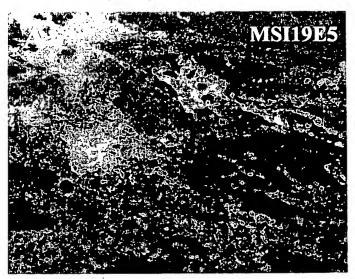


FIG. 27B

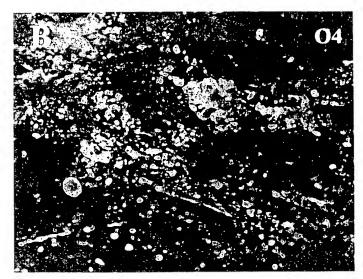
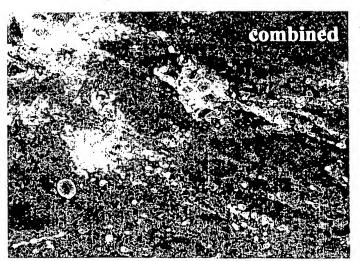
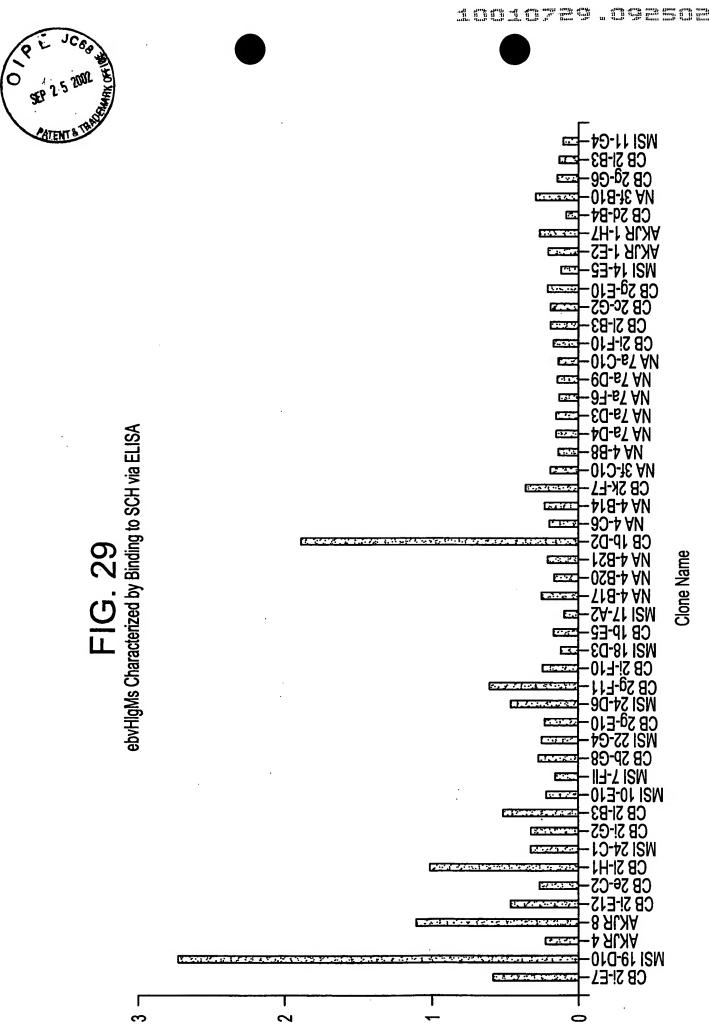


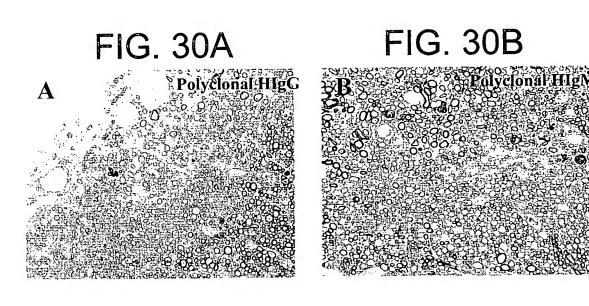
FIG. 27C

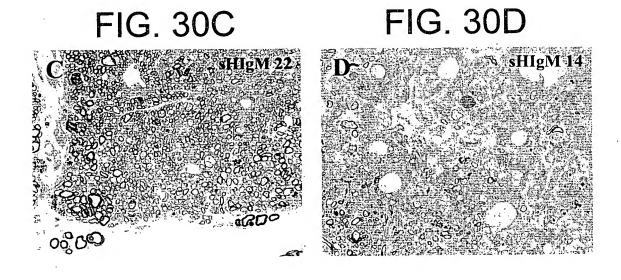


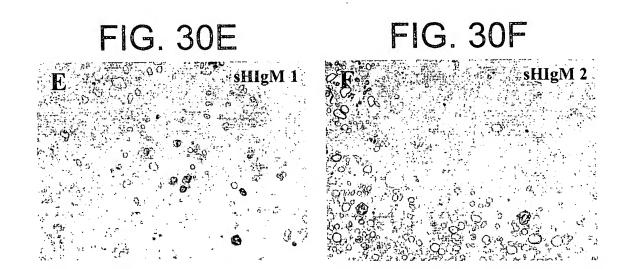


(mn cu+) .u.U













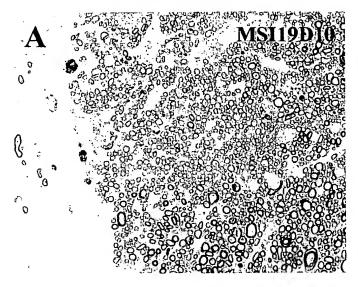
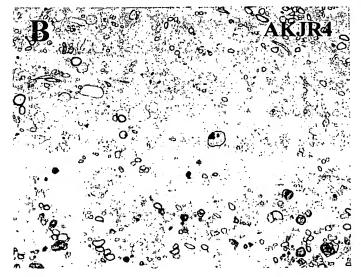


FIG. 31B







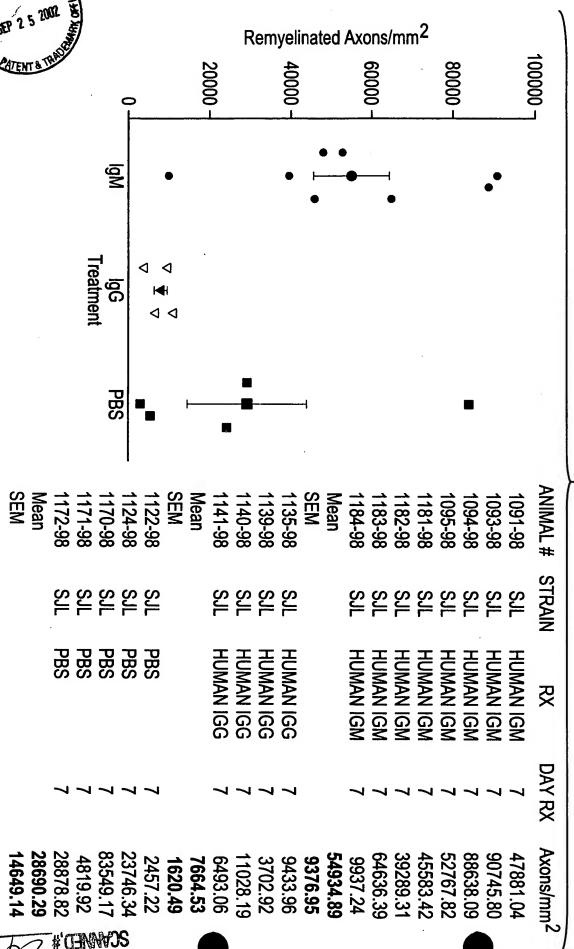
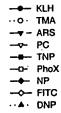
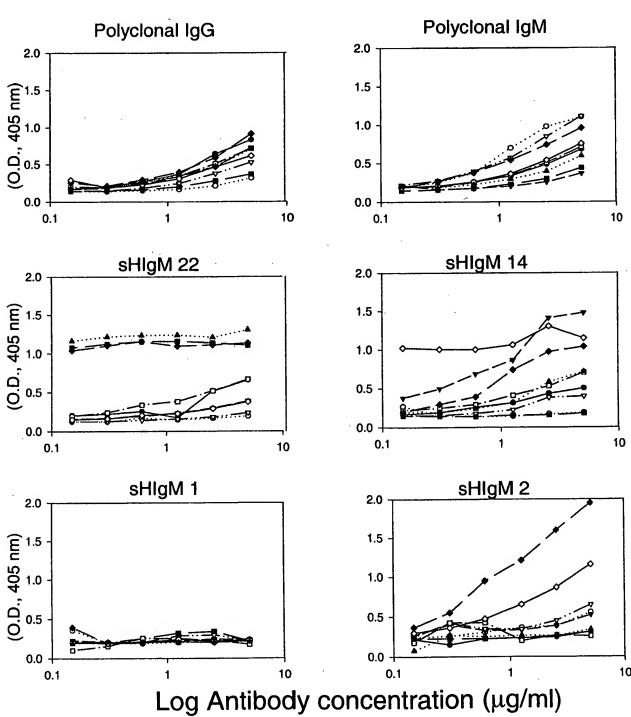




FIG. 33

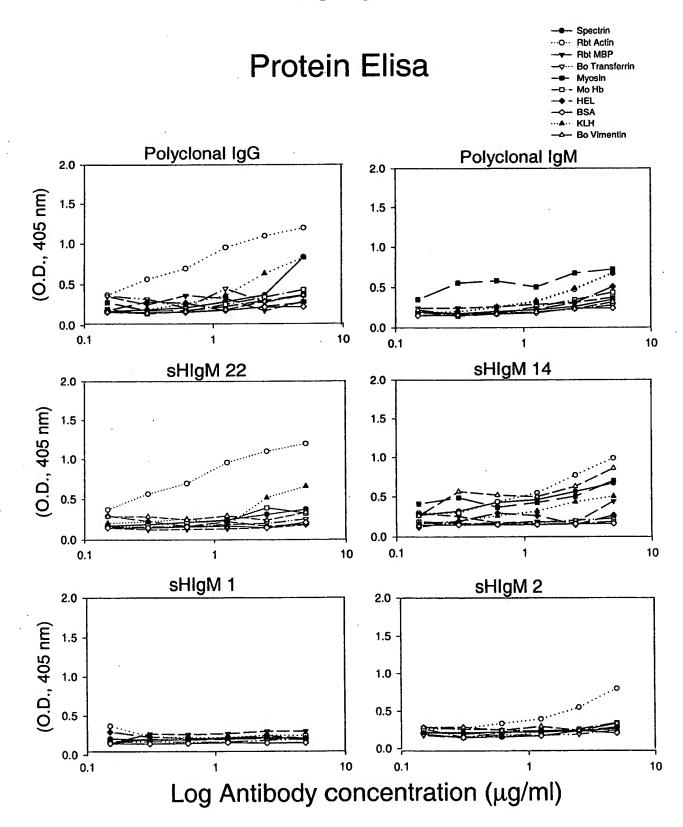
Hapten Elisa













/FR	1													
1	2						8 <i>G</i>						14 P	
CAG	CTG	CAG	CTYG	V GTG	GAG	TCT	GGG	GGA	GGC		TC GTC			<u>G</u> GGG
	e A sl				0.10		-	G						
	e B sl	_												
16	17	18	19	20	21	22	23	24	25 .	26	27	28		
R	<u></u>	L	R	L	<u>S</u>	_ <u>C</u> _	A	<u>A</u>	S	$\frac{G}{G}$	<u>F</u>	T	F	S
AGG	TCC	CTG	AGA	CTC	TCC	TGT	GCA	GCC	TCT	GGA	TTC	ACC	TIC	. AGI
/CDI	R1				-/FR2	?								
31	32	33	34	35			38							
S	S	G	M	H	W		R	Q	A					
AGC		GGC	ATG	CAC	TGG	GTC	CGC	CAG	GCT	CCA	GGC	AAG	GGG	CTG
	C													
			,	/ CDR	2									
46	47		49	50	51		52A					57	58	59
E	W	V	A	V(I	<u>I</u>	S	Y_	D	G_	S	R	K		
GAG	TGG	GTG	GCA	GTT		TCA	TAT	GAT	GGA	AGT		AAA	TAC	TAT
				A C	T T						GG GG			•
						/FR3								
60	61	62	63	64	65	66	67	68	69	70	71	72	73	74
<u>A</u>	D	S	V	K	G	R	F	T	I	S		D	N	<u>_S</u>
GCA	GAC	TCC	GTG	AAG	GGC	CGA	TTC	ACC	ATC	TCC	AGA	GAC		TCC
													C	
											·			
75	76	. 77	78	79	80	81	82	82A	82B	82C	83	84	85	86
K	N	T	L	<u>Y</u>	L	0	M	N	S	L	T	<u>A</u>		<u>)</u> D
AAG	AAC	ACG	CTG	TAT	CTG	CAA	ATG	AAC	AGC	CTG		GCT	GAG	GAC
		T	~								CG C		C	
		T	C 					/ CDR	3					
87	88	89	90	91	92						98	99	100	100A
T	A	V	Y	Y	C	_	K	_		T	G	S	P	T
ACG	GCT	GTG	TAT	TAC	TGT	GCG	AAA	GAG	GTG	ACT	GCT	ATT	CCC	TAC
				T				GA			G	G		ACG
			1004					GA			G	G	G	ACG
100	D101		/FR4		105	106	107	108	100	110	111	112	113	•
							T							
$\overline{\mathrm{TTT}}$	GAC	TAC	TGG	GGC	CAG	GGA	ACC	CTG	GTC	ACC	GTC	TCC	TCA	
C										•			G	;
С													G	;
•														



/FRI						_								
1												14		
0	S		L	_T_	_0_	P	<u> P</u>	S		S	_A	_A_	P	G
								TCA	GTG	TCT	GCG	GCC	CCA	GGA
		l-IgM.					${f T}$				T			
Clon	e II sł	H-IgM	.22 V	λ	G		T				${f T}$			
						/	CDR1	L						
		19	20	21	22	23	24	25	26	27	27A	27B	28	
<u>Q</u>	<u>K</u>	<u></u>	<u>T</u>	<u>I</u>	_ <i>S</i>	_ <u>C</u> _	<u>_S_</u>	_G_	<u> </u>	_ <u>S_</u>		N	<u>I</u> _	_ <u>G</u>
CAG	AAG	GTC	ACC	ATC	TCC	TGC	TCT	GGA	AGC	AGC	TCC	AAC	ATT	
														C
					/ mm 2									<u> </u>
30	21	32	22	/ 21	7 F R Z =	36	37	30	30	40	A 1	42	43	11
												T		
<u> </u>	AAT	TAT	GTA	TCC	TGG	TAC	CAG	CAG	СТС	CCA	GGA	ACA	GCC	CCC
		T	0111	-00			0.10	A	0.0	00				
		T						A						
					CDR	2						-/FR3	3	
												57		
												G		
	CTC	CTC	ATT	TAT	GAC			AAG	CGA	CCC	TCA	GGG	ATT	CCT
G						T T	C							
						T								
60	61	62	63	64	65	66	67	68	69	70	71	72	73	74
D		F				K								
												ACC		
•														
75	76											87		
	T	G	L		T	G	D	E	A	D	<u>Y</u>	<u> Y</u>	<u>C</u>	G(E)
ATC	ACC	GGA	CTC	CAG	ACT	GGG	GAC	GAG	GCC	GAT	TAT	TAC	TGC	GGA
														_
										/ 1710 A				A
	01			0.4	0 E	053	OED			/ FR4		100	101	102
90 70	9 T	92	93	94	95 T.	93A	93B 8	70 17	3 / 17	JO F	23 C		101 G	TOZ
7 7	TCC	GAT.	AGC	AGC	ርጥር		 ጥ	GTG	GTA	ጥጥር	GGC	GGA	GGG	ACC
ACA	133	ONI	AGC	AGC	C10	AGT	GC	010	OIM			G		
						AGT						G		
						/ Cλ			_			_		
103	104	105												
		T												
AAG	CTG	ACC	GTC	CTA	GGT	CAG	CCC	AAG						



Sequence of MSI 19-D10 VH

FR1															
1 CAG		3 CAG	CTG	CAG	GAG	TCG	GGC	CCA	GGA	CTG	GTG	AAG	CCT	15 TCG S	
														/ CDR	1
ACC	18 CTG L	TCC	CTC	ACC	22 TGC	23 ACT	24 GTC	25 TCT	26 GGT	27 GGC	28 TCC	29 ATC	30 AGT	31 AGT	,
		- -	,	FR2											
32 TAC	33 TAC Y	34 TGG	35 AGC	36 TGG	37 ATC	38 CGG	39 CAG	40 CCC	41 CCA	42 GGG	43 AAG	44 GGA	45 CTG	GAG	
			CDR	12											
47 TGG	48 ATT I	49 GGG	50 TAT	51 ATC	52 TAT	53 TAC	54 AGT	55 GGG	56 AGC	57 ACC	58 AAC	59 TAC	60 AAC	61 CCC	
				FR3											
TCC	63 CTC L	64 AAG	65 AGT	66 CGA	67 GTC	68 ACC	69 ATA	70 TCA	GTA	GAC	ACG	TCC	AAG	AAC	
CAG Q	78 TTC F	TCC S	CTG L	AAG K	CTG L	AGC S	TCT S	GTG V	ACC T	GCT A	GCG A	GAC D	ACG	GCC	-
															_
GTG	90 TAT Y	TAC	TGT	GCG	AGG	TCG	GCA	CAG	CAG	CAG	CTG	GTA	TAC	TAC	:
		/	FR4											/Cµ-	
100I TTT	0 101 GAC D	. 102 TAC	2 103 TGG	3 104 GGC	105 CAG	5 106 GGA	5 107 ACC	7 108 CTG	3 109 GTC	9 110 ACC) 111 GTC	L 112 TCC	2 113 TCA	3 114 GGG	



Sequence of MSI 19-D10 $V\kappa$

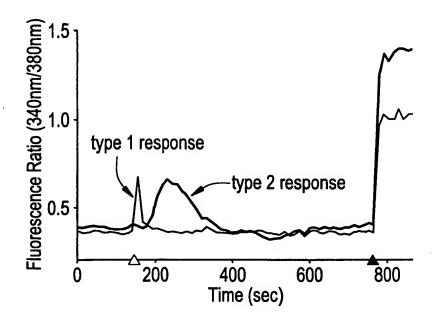
FR 1														
1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
GAC	ATC	GTG	ATG	ACC	CAG	TCT	CCA	GAC	TCC	CTG	GCT	GTG	TCT	CTG
D	I	V	M	T	Q	S	P	D	S	L	A	V	S	L
								, CDE)1					
16	17	18			21									
	GAG	AGG	GCC	ACC	ATC	AAC	TGC	AAG	TCC	AGC	CAG	ACT	CTT T	2 / C ጥጥል
G	E	R	A	T	ī	N	C	K	s	s	0	S	v	L
_				_	_	-•				_	-	_	•	_
		~~~								/FR2				
	27E													
TAC	AGC S	TCC	AAC	AA'I'	AAG	AAC	TAC	TTA	GCT	TGG	TAC	CAG	CAG	
1	3	3	7.4	N	K	14	I	ם	A	W	I	Q	Q	
											٠			
										/	CDR	2		
	40													
	CCA													
K	P	G	Q	P	P	K	L	L	I	Y	W	A	S	T
			FR3											
	55												67	68
	GAA													
R	E	S	G	v	P	D	R	F	S	G	S	G	S	G
69	70	71	72	73	74	75	76	77	78	79	80	81	82	83
	GAT													
	D													
														. == 4
					/	_								/FR4
	85													
	GTT													
A	V	Y	Y	C	Q	Q	I	I	5	T	P	L	T	F
									/Cĸ					
99					104			-			110	111	112	113
	CCT													
~	73	~	m	77	77	~	_	TP	*	70%	~~	-	-	~



### **FIG. 39A**

Mixed Primary Glia sH-lgM.22 Ca²⁺ response

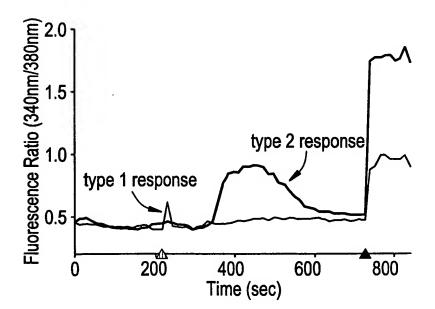
- ratio cell #1
- ratio cell #2
- $\triangle$  sH-lgM.22 (3 $\mu$ g/ml)
- ▲ Br-A23187 (10μM)



### FIG. 39B

Mixed Primary Glia SCH 94.03 Ca²⁺ response

- ratio cell #1
- ratio cell #2
- Δ SCH 94.03 (3µg/ml)
- ▲ Br-A23187 (10μM)



### FIG. 39C

Mixed Primary Glia
CH 12/sH-lgM.14 Ca²⁺ response

- ratio cell #1
- ratio cell #2
- A CH 12 (3μg/ml)
- $\triangle$  sH-lgM.14 (3 $\mu$ g/ml)
- **A** Br-A23187 (10μM)

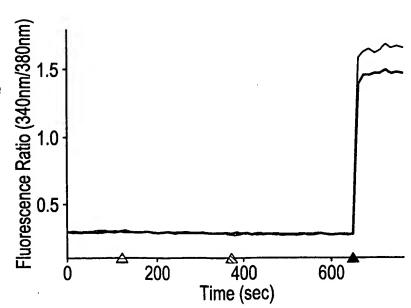




FIG. 40A

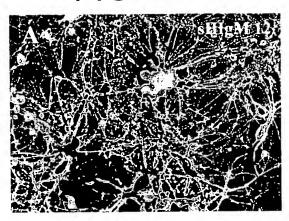


FIG. 40B

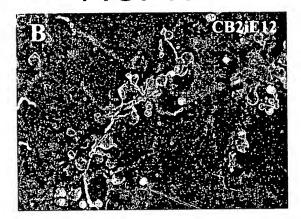
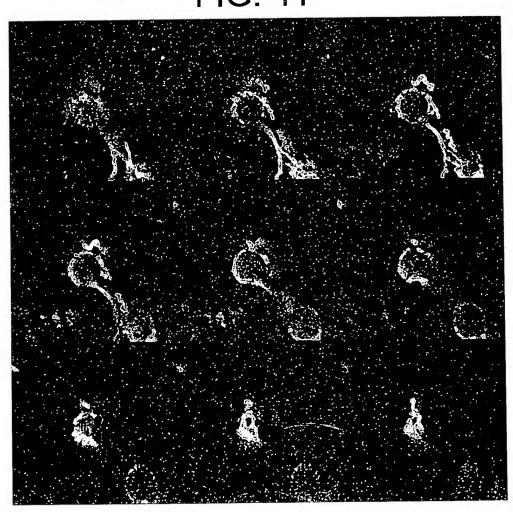




FIG. 41







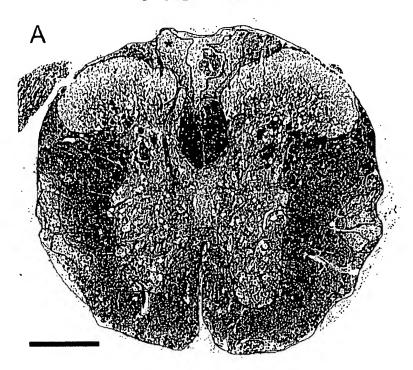
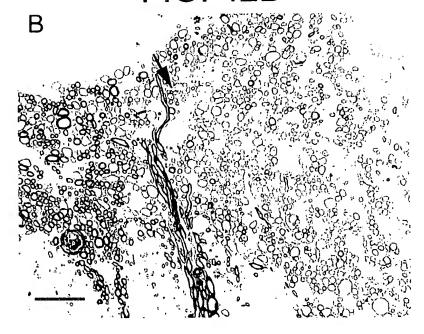


FIG. 42B





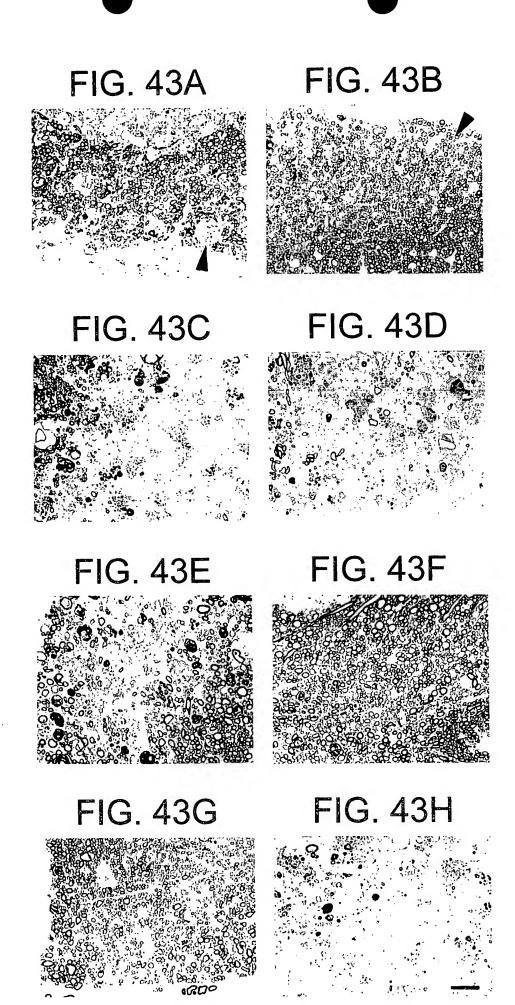


FIG. 44A

FIG. 44B

FIG. 44B

FIG. 44B

FIG. 44B

### Translation of CB2b-G8 $V_{\rm g}$ :

1				5				x	10	A	v	v	0	15 P	G	R	S TC	L	20 R	L	
C TGT	A GCA	25 A GCG	S TCT	G GGA	F TTC	I ATT	30 F TTC	S AGT	S AGC	R1 - Y TAT	IMG7 G GGC	7 35	•••	• • •	•••	< M ATG	40 H CAC	W TGG	V GTC	R CGC	Q CAG
45 V	P	G	к	G	50 L	E	W	V GTG	A	55 V	1	W TGG	Y TAT	CDR2 D GAT	2 ~ ; 60 G GGA	IMGT S AGT	D GAT	K AAA	•••	65	Y TAC
																	I K AAA				Y TAT
L	90 Q CAA	м	N	s	L	95 R	A	E	D	т	Α	v	Y	Y	С	105 A	R AGA	D	R	S	S
											G GGA						125 S TCC	S TCA			



### Translation of CB2b-G8 $V_{\lambda}$

< F	R 1 - I M G T	
1 5	10	15 20
× 	X L L S G S  XGC CTC CTG TCT GGG TCT	P G Q S I T I S CCT GGA CAG TCG ATC ACC ATC TCC
		·
	CDR1 - IMGT	<b>&lt;</b> -
25 C T G T S S D	30 CDR1 - IMGT 35 V G G Y N Y	40 V S W Y O O
CTG ACT GGA ACC AGC AGT GAC	GTT GGT GGT TAT AAC TAT	GTC TCC TGG TAC CAA CAG
	_	_
		CDR2 - IMGT65
45 50 H P G K A P K	L M I Y D V S	60 65 D
CAC CCA GGC AAA GCC CCC AAA	CTC ATG ATT TAT GAT GTC AGT	D GAT
	_	
•	F	
70 75 R R S G V S	80 N R F S G S K	85 SGNTAS
CGG CCC TCA GGG GTT TCT	AAT CGC TTC TCT GGC TCC AAG	TCT GGC AAC ACG GCC TCC
		>
		CDR3 - IMGT
LTISGLO	100 A E D E A D Y	YCSSYTSS
CTG ACC ATC TCT GGG CTC CAG	GCT GAG GAC GAG GCT GAT TAT	TAC TGC AGC TCA TAT ACA AGC AGC
	115 120	125
S S V V F G G	G T K L T V L	G Q P K A A P S
AGC TCT GTG GTA TTC GGC GGA	GGG ACC AAG CTG ACC GTC CTA	GGT CAG CCC AAG GCT GCC CCC TCG

FIG. 47A

DHFR amplification of 94.03k 2

0.2 51.2 Neg Pos 0.2 51.2

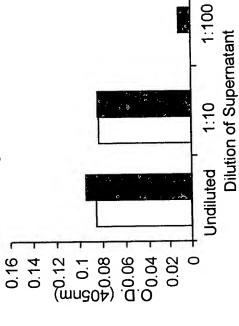
### FIG. 47C

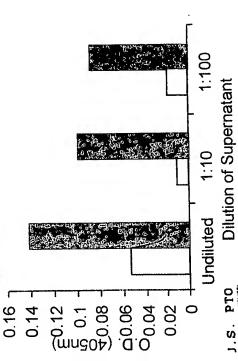
Clone #5 Kappa Chain Elisa

0.2 ug/ml methotrexate 51.2 ug/ml methotrexate

0.2 ug/ml methotrexate 51.2 ug/ml methotrexate

Clone #4 Kappa Chain Elisa





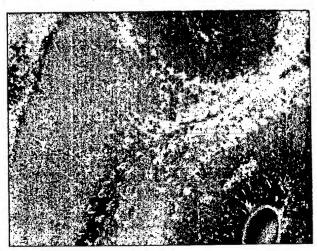






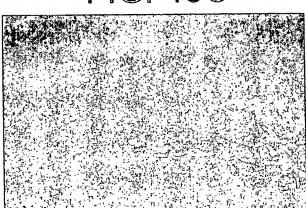
Mouse 94.03

FIG. 49B

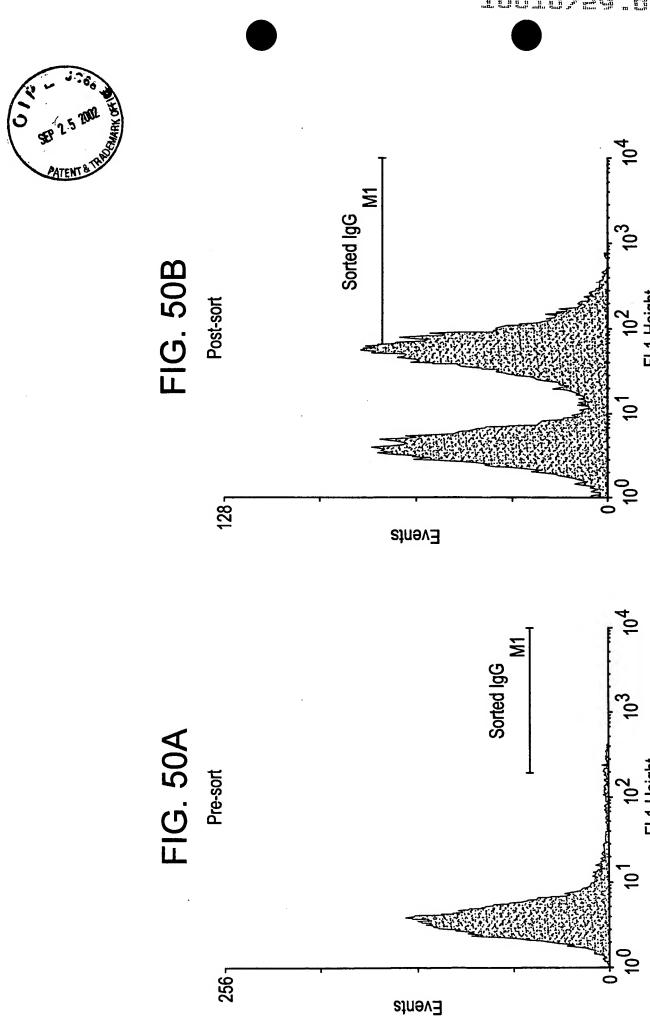


Humanized 94.03 clone 1

FIG. 49C

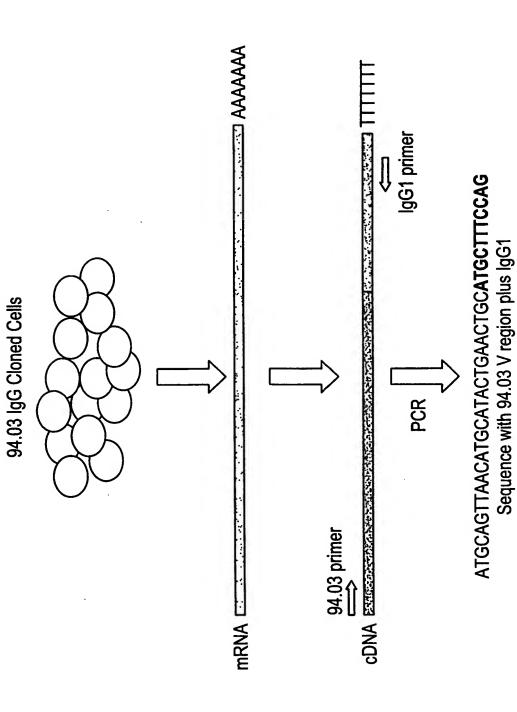


Humanized 94.03 clone 2





**FIG. 51** Sequencing of 94.03 lgG





### 09 $V_{\scriptscriptstyle R}$ Sequence with translation:

<						F	R	1	-	I M	G	т									
1			Ę	5				1	١٥				1	5				2	0		
Q	D	Н	L	0	Q	S	G	P		E	L	V J	K	P	G	Α	F	V			
CAG	GAT	CAC	CTG	CAG	CAG	TCT	GGA	CCT		SAG (	CTG (	STG F	AAG	CCT	فاتات	GCT	TTT	GTG	AAG	ATA	TCC
			>													<					- <b>-</b>
					25				CDR	1 30	imgt				35					40	
С	K	A	S	G	Y	T	F	T	N	Y	D GAT					L	N	W	V	R	Q
TGC	AAG	GCT	TCT	GGT	TAC	ACC	TTC	ACA	AAC	TAC	GAT	• • •	• • •	• • •	• • •	CTA	AAC	166	GIG	AGG	CAG
F	R	2	-	I	M	G	T ·			>											<
-														CDR2	- I	MGT					
45					50	_		_	_	55	I ATT			_	60	-				65	v
R AGG	P CCT	GGA	Q CAG	G	L CTT	E GAG	W TGG	I ATT	G GGA	W TGG	ATT	Y TAT	CCT	GGA	N AAT	GAT	N AAT	ACT			. AAG
		00	00	-																	
												-	F	R	3	-	I	M	G	Т	
							_					_						or			
Y	N	E	70 K	F	ĸ		G 7	75 L	A	s	L	Т	A A	D	ĸ	s	s	85 T	T	A	
TAC	AAT	GAG	AAG	TTC	AAG	0	GGC (	CTG	GCC '	TCA	CTG .	ACT (	GCA	GAC A	AAG	TCC	TCC	ACC	ACA	GCC	TAC
															>						
																					110
τ.	90 H	т.	g	g	T.	95 T	s	E	S	s	100 A	v	¥	F	С	105 A	R	G	L	Р	R
TTG	CAT	CTC	AGC	AGC	CTG	ACT	TCT	GAG	AGC	TCT	GCA	GTC	TAT	TTC	TGT	GCA	AGA	GGC	TT	A CC'	r agg
ann		T) 40 E													•						
CDR.	<b>5</b> - 1	IMGT		115		_		12	0												
G	W	Y	F	D	v	W	G	Α	G	T	T	V	T	V	S	S	A	,			
GGC	TGG	TAC	TTC	GAT	GTC	TGG	GGC	GCA	فافاقا	ACC	ACG	GTC	MUC	. GIC	100		. GCI	•			



# Translation of 09 kappa light chain 1:

		ACC .
		r Trg
•	!	20 T ACC
		20 V T GTC AC
••	1 1 1	R AGG
ဌ	1	E GAG
chain 1	1	ი გეგ
	: : : : :	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$
light	€	S TCA
•	ליז	M ATG
карра	Σ	S
0 0	н	M C ATG T
) H O	ı	10 TCC
ŭ O	Н	AAA
1410	ద	4 222
Translation	Ĺ	S
Irar	 	CAA
<b>5</b> 1		5 ACC
3		M ATG
53	)	I V M ATT GTA ATG
(		ATT
ū	• ;	1 AAC

									ı			K.
		ø		>			z	AAC	۲		ſĿι	TTC
1		Ø				65		:	ტ		Ω	GAT
			TAT					:	Ħ	85	₽	ACA
			TGG					:	Н		ď	gC _A
	40	ഗ	r TCC					:	1		ഗ	ICI
i V	ı	>	. GI		IMGT			:	m			:
			:		1	9		:	r <del>c</del>			:
			:		CDR2			: ບ	Įτι	80	ტ	GGA
			:				S				ഗ	
	35		:				ď		1		ტ	
IMGT			:				ტ				E٠	
1			:	?		55	Þ	Ŧ	į			
CDR1		¥	TAT	1			Н	ATA	1		Įτί	
Ü			ACT				L	CTG		75	硂	
	30		GTT	Ŀ			ᄓ		1		Д	GAJ
	ı		GTG	۔ ن			×		!			:
		z	AT	) M		20	ሲ				щ	CCC
		ធ	GAG A	_			ഗ				>	GTC
<b>?</b>		ß	AGT (				o		1	70	ტ	999
	25		₹ 225 5	7			回	3AG (	1 1		E	ACT
i !	. 4	×	AAG (	R			М	SCA (	1		×	TAC
1		ပ	TGC 7	L		45	×	AAA CCA GAG (	1			CGG

	<u>-</u>
	IMGI
	۳
	CDR
	Ì.
<b>^</b>	
!	
į	
į	
į	
	•
1	
1	
1	
- 1	
-	
1	
! !	

S A T D F T TCT GCA ACA GAT TTC ACT

110	×	TAT
	တ	AGC
	×	TAC
	ტ	GGT
	Ø	CAG
105	ტ	GGA
	ပ	TGT
	Ħ	CAC
	×	TAT
		GAT
100		GCA
	ᄓ	CTT
	Ω	GAC
	ы	GAA
	ď	GCT
95	Ø	CAG
	>	GTG
	ഗ	AGT
	ഗ	AGC
	Н	ATC
90	H	ACC
	Ы	CTG

P Y T F G G GC CCG TAC ACG TTC GGA GGG GGG



Translation of 09 kappa light chain 2: FIG. 54

				1	:
	n AAT	GAG	ACT	TACT	IMGT 110 3 Y AA TAC
	I ATT	O.A.	65	F F TTC	E GAA
	20 T ACT	Y		GAT	CDR3 N AAT
	I ATT	TGG		85 ACA	H
 	T ACC	40 A		н 667	Q CAG
	E GAA	 L TTA	IMGT	s TCT	105 Q CAA
1	ი გეგ		H :	m :	TGT L
	15 P CCT	. :	CDR2	<u>د</u> :	TAC
€+	F TTT	:	l so	80 80 66A	Y TAT
ტ	[독] ^S	35	o gg A	S AGT	M ATG
Σ	A GCT	IMGT	S TCT	၂ ဗဗ္ဗ	100 A GCA
н	CTT	H + :	55 TAC 1	S	F
1	10 Y TAT	CDR1 Y TAT	ATC	FF	DGAT
-	STCT	A A A A	110	75 R AGG	GAA
ø	CCA ,	30 S AGT	T L CTT (	S TCA	CCT TCCT
[t4	S ICT (	I ATT 2	G K A A A G C	:	95 BE GAG
!	O CAG	S AGC 2	M (50 N AAT 1	GCA CCA	r. CIG
	T ACC (	K AAG	H H H	IATT	S AGC
į	1 1	S S AGT 2	K AAA	70 G	SAGT
!	V Q I GTC CAG ATA	25 A GCA 7	2	s ,	ATC
į	V FTC (		R 2	CAA	90 T ACC 2
>	D GAT C	C R TGT AGG	F F F R R AGA C	TTG (	CTC

115 P Y T F G G G CCG TAT ACG TTC GGA GGG



## Translation of AKJR 4 Heavy Chain:

FIG. 55

1	<b>7</b> )	1 (1)		د	:	E	o v
į	s TCC	CAG	i 1	Y TAT	į	F TTT	MGT 11. CC
į !	r CTC	, a C			۲	v GTG	- IMGT 110 G P GGT CCC
	20 R AGA	V GTC		:	v	T ACG	CDR3
	CTG	™ TGG		rg s	Σ	85 S AGC	E
] ] 	s TCC	40 S AGC		SAGT	н	K AAG	CAG
	ა ე	M ATG		· -	ı	S	105 A GCG
į	999 9		- TMGT	E		NAAT	TGT
•	15 P CCT 0				œ	D GAC 1	TAC
1			£	, _			! _
Ħ	CAG	:		ြည်	Œ	80 R AGA	TAT
ტ	STA	35		SAGT	,	STCC	I
E	TIG	Gr GCC		Ciri		IATC	100 A GCC
н	ဗဗ	Y A	Ŷ	55 S AGT	į	T ACC	ACG
ı	10	CDR1 D	1	s TCA		F	GAC
	g GGA	ATC		v GTC		75 R : CGA	E GAG
æ	უ ე	30 F	ا 1	¥ TGG		6 66C	<b>₽</b>
[24	S TCT	S AGC	ტ	E GAG		•	95 R AGA
į	E GAA	F	Œ	50 L CTG	 	K AAG	r. CrG
1	S L TTG (	GGA '	н	g G		V GTG	S
	L CITA	S TCT (		K AAG	!	, D	S
i !	CAA	25 A GCC :		9		70 D S GAC TC	L CTG
!	v GTG	A GCA G	<b>8</b>	ч 2)	į	A GCA O	90 08 08 08
>	1 E GAG G	C C	E4	45 A GCT C	1	Y TAT	CTG C
•							

Q R R W G Q G T L V T V S S G S A S A P T L CAG CGT CGC TG GGA ACC CTG GTC ACC GTC TCC TCA GGG AGT GCA TCC GCC CCA ACC CTT



### FIG. 56 Tran

# Translation of AKJR 4 Kappa Light Chain:

	T ACT	CAG		N AAT		T ACT	IMGT 110 Y T TAC	
1	IATC	CAC	٧	65	E	F	- IM S AGT	
	20 T ACC	Y TAT		:	Ŋ	E GAA	CDR3	
1	V GTC	TGG		:	Σ	85 T ACA	Y TAT	
·	R AGA	40 A DDG		:	н	8 5	CAG	
!	DGAC	L ITIG	Ę		ŧ	S	105 Q CAG	
!	ဗ ဗဗ္ဗ	1	Ì		м		ဂို ပဋ	
; ; ;	15 V GTA		Ç		æ	•	Y	
E	STCT		į	F. T.T.	ĵz,	0 0 0 0 0 0	Y	
O	A GCA	35		<b>₽</b>		S S AGT G	T. ACT.	
×	STCT			K AAG		ပ ပ	100 A GCA	
н	CTG	- IMGT	^	y TAT	1	R	S	
ı	10 T ACC	CDR1		I ATC		F	D	
7	S TCC	S AGC		r CTG		75 R AGG	D	
ĸ	PCCT	30 S AGT	ا 1	r Crc		s TCA	PCCT	
Œ	S TCT	IATT	ტ	K AAA		•	95 O CAG	
	CAG	SAGT	×	50 P CCT	!	CCA P	r CIIG	
   	5 T ACC	CAC	н	<b>A</b>		V GTC	S	
	MATG	S AGT	ı	K AAA	1 1 1	70 G	SAGC	
!!!	Q CAG	25 A GCC	7	ა მ		S SAGT	I ATC	
 	IATC	R CGG	<b>6</b> 4	P CCA		GAA	90 T ACC	
     	1 D GAC	ngc 1	ſΞι	45 K AAA	! ! !	r TTA	CTC	



## Translation of CB2i-E12 Heavy Chain: FIG. 57

.	s TCC	O CAG	AAC	Y TAC	110 Y TAT
	V GTC	R CGA	65	T A GCC	S
	20 K AAG	V		G ACA	R CGA
	V GTG	TGG	T. ACA	M 85 S AGC	D
	S TCA	40 H CAC	5 5 5	I I ATC	R AGA
1	A GCC	M ATG	IMGT _ ) G G ST GGT	r S T	105 A GCG
.	E GAG	:	- IN 60 S AGT	w F1 D V	To T
; ; ;	15 x XCG		CDR2 N	R GAC	Y TAC
E	K AAA		D P CCT	F R AGG	Y TAT
O	× AXG	35	N	ACC	V GTG
Ħ	× XAX	IMGT _	I	MATG	100 A GCC
н	× XAG	- IM Y TAC	-> 55 W TGG	ACC	T ACG
ı	10	CDR1 G	G	V GTC	D GAC
႕	r Agg	C L	M ATG	75 R AGG	D GAC
æ	× S	30 F	T - W TGG	၁ <del>၅</del>	S TCT
្រុ	•	T	G E GAG		95 R AGA
<u> </u>	•	Y	M 50 L CTT	O CAG	L CTG
 	٠ .	GGA	ы В В	F. T.T.T.	R AGG
i !	· •	S S TOTE	CAA	70 K AAG	S
; ! !	•	25 A GCT	S GGA GGA	OCAG	r r
 	:	K K AAG (	A GCT	A GCA	90 E GAG
>	ਜ :	2 U	F 1 45 A A GCC (	Y TAT	MATG

P G R N Y F D Y W G Q G T L V T CCG GGA AGG AAC TAC TTT GAC TAC TGG GGC CAG GGA ACC CTG GTC ACC CDR3 - IMGT



## Translation of CB2i-E12 kappa chain: FIG. 58

	S	Q CAG	   	AGC	T ACT	IMGT 110 S TCT
	CIC	Q CAG	65	: €	F TTC	- S AGC
	20 T ACC	Y		: o	D	CDR3
	A GCC	W TGG		: ¤	85 T ACA	Y TAT
!	R AGA	40 A GCC 1		: н	Ŋ	O CAG
 	E GAA	L TTA	IMGT _	: 1	S TCT	105 0 CAG (
i !	9 9		МІ - 09	. m	. : '	C C TGT
i !	15 P CCA (	:	CDR2	: £	:	Y
E+	s TCT		) '	သင်္ဂ	80 9 9	Y TAT
Ŋ	L TTG'	3.5		A GCA	8 S AGT	V GTG
×	S TCT	GT		GGT	9 9	100 A GCA (
н	L CTG	- IMGT Y TAC	V 555	TAT	S AGT	TTT
1	10 T ACC	CDR1		ATC	F	D
н	၁၅၅	- S		CHC	75 R ? AGG	E GAA
ρζ	P CCA	30 S AGC 7	E+ '	GFC	D GAC	CCT
Įτι	S TCT	V GTT	<b>o</b>	AGG	•	95 E GAG
	O CAG	S	M 50	CCC	CCA	CTG
1	5 T ACG	Q CAG	н	GCT	IATC	R
İ	L TTG	S AGT		CAG	70 G G G G	S
! ! !	V GTG	25 A GCC 7	0	၁၅၅	70 T G ACT GGC	I
; † 	I ATT (	R AGG 0		CCT	A GCC	90 T
· •	1 E GAA 2	0 1 0 1		AAA	R AGG	L

¹¹⁵ H T F G Q G



## Translation of CB2i-E7 Heavy Chain: FIG. 59

) ) 	S		Q CAG	;		N AAC	1 1 1	Y TAT	
1 1	CHC		R GGC	٧	65	:	E	L CTG	
1    - 	20 R AGA	1	IATC			:	ტ	S TCA	
 	CTG		TGG.			T ACA	×	85 N AAC	
	S	40	S AGC			Y	н	K AAG	
 	999 9	1	MATG	E-	1	S AGT	1	A GCC	
1	g GGA		:	Ž.	!	SAGT	m	N AAC	^ ¹
	15 P CCT (		•	CDR2		SAGT	ద	D GAC	
Ē	K AAG (		:	5	j	S AGT	[24	80 R AGA (	
Ŋ	V GTC	35	:		ŧ	SAGT		နှင့် မရှင်	1
Σ	T.T.G	IMGT	Y TAC		j.	IATT	! ! !	I ATC	
н	ဗ္ဗဗ္ဗ	1	Y TAC	<b>^</b>	55	Y		TACC	
1	10	CDR1	D GAC	     		S	1 1	F	
-	× &		S AGT			V GTT	.	75 R CGA	
<b>~</b>	:	30	F	ı E		₩ TGG	 	ဗ္ဗ	
Ľτι	:		T ACC	ტ		E	1	:	1
1	:		F	Σ	20	L	ļ	K AAG	
!	ري :		G GGA	н		ი მ		V GTG	
	:	<b>^</b>	S TCT	1		K AAG	! ! ;	70 s TCT	
	:	25	A GCC	7		G GGG		D GAC	 
!	•		A GCA	æ		CCA		A GCA	
\ \ \	н :	!	CTGT	[t4		A GCT		Y	1

110	ഗ	AGC	
	ഗ	TCG	
	æ	SSS	
	Д	GAT	
	ፚ	AGA	
105	¥	909	
	ပ	$\mathtt{TGT}$	
	×	TAC	
	≻	TAT	
	>	GTG	
100	Ø	GCT	
	H	ACG	
	Ω	GAC	
	ធា	GAG	
	Ø	ပ္သင္ဟ	
95	吆	AGA	
	ü	CTG	
	ß	AGC	
	z	AAC	
	Ħ	ATG	
90	Ø	CAA	
	ı	CTG	

S S W Y Y Y G M D V W G Q G AGC AGC TGG TAC TAC TAC GGT ATG GAC GTC TGG GGC CAA GGG CDR3 - IMGT



## Translation of CB2i-E7 kappa Chain: FIG. 60

ı					ŧ		. 0
!	T ACT	, O &		E	ACT	T ACT	IMGT 110 C TGC
	IATC	0 8	Ť	65	E	F TTC	- K K AAG
	20 T ACC	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	4		: ₀	D GAT	CDR3
1	V GTC	] 3 E	3		<u> </u>	85 T ACA	Y
	r Aga	40 A A	3		: :	9 9	K
!	DGAC			 	: 1	S TCT	105 0 CAA 2
1	GGA .		·	- TW	: m	: :	C C TIGIT
1	15 V GTA				; ¤	: 3	Y TAC
H	S TCT		: :	ر ا ه	TCC F	80 G GGA	Y
ტ	A GCA	35	• •	Æ	GCA	S AGT	T
×	STCT	IMGT -	• •	A	GCT	g 0	100 A GCA 7
н	r CTG	Σ H I	: ^	55 Y	TAT	n AAT	V GT'T
ı	10 s TCC	CDR1		н	ATC	F	D
ᡤ	s TCC	X 4		ᄓ	o I o	75 R CGG	E
ĸ	P	080 080	· E	13	o l	S	P
Ľι	s TCT	H	U	×	AAG	:	95 O CAA
	Q CAG	ဗ ပွ	) ) )	50 P	CCT	CCA	L
!	5 T ACC	0 8		>		V GTC	S
	M ATG	\ - \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \		×		7.0 G GGG	S
! !	Q CAG	25 A A		တ		S TCA	I
.   .   .	IATC	M C		Д		o Caa	90 T ACC
  -   	1 D GAC			45 K	1	L	CTC

P S H F R G R D CC TCT CAC TTT CGG GGG AGG GAC



Translation Of MSI 19-E5 Light Chain

1 D I A M T Q S P D S L A V S L G E R A T I N GAC ATC GCG ATG ACC CAG TCT CCT GCA GTG TCT CTG GGC GAG AGG GCC ACC ATC AAC ט Σ н

		ø	CAG	· · · · · · · · · · · · · · · · · · ·		E	ACC	! !		H	ACT		Η -	110	ACT
		Q	CAG		65		:	€÷		Ŀ	TTC		CDR3		S AGT
!		×	TAC		:		:	ტ		Ω	GAT				Y TAT
1		Z	TGG				:	Σ	85	E	ACA			:	Y TAT
1	40	æ	GCT				:	н		U	999				G &
		ы	TTA	IMGT			:	1		ഗ	ICI			105	CAG C
		þ	TAC		09		:	ო			:	^ !		,	r TGT
		z	AAC	CDR2			:	œ			:			:	TAC
		z	AAC			Ø	TCT	Ē	80	U	999			;	Y TAT
	35	z	AAT			Ø		,			AGC	1			V GTT
IMGT		z	AAC			3		1		ტ	GGC	· I		10	4 25 42
Ξ.		ഗ	ICC	ì	. 55	Þ	TAC			ഗ	AGT	1			V GTG
CDR1		ഗ	AGC			н	ATT	1			TIC				D GAT
·		Ēų	TTC			ы	CIC		75		CGA	1			GAA
	30	ы	TTA	Ħ		ы	CIA			Ω	GAC	1			GCT
		>	GTT	υ		×	AAG				:			95	વ કુ
		ഗ	AGT	Σ	50	Д	CCT			Д	CCI	į			L CTG
		ద	CGG	н			CCT	i, i			GTC	i 1			S AGC
<u>^</u>	•		AGC	1		Ø	CAG	1			999	1			S AGC
	25			7		ტ	GGA				TCC				IATC
				æ			S	1			GAA		 	90	T ACC
i !		υ	TGC	ţzı	45	×	AAA	!		œ	CGG	1	 		CIC

MGT

P I T F G CCA ATC ACC TTC GGC



### Translation of 04 kappa chain 2: FIG. 62

	T ACC	OCAG	 	Y	   	T ACT	IMGT 110 T T CT ACT
1	I ATC	40 Q CAA	<b>V</b>	:	E	F	- II
!	20 S AGC	 4 Y TAT		:	ဗ	D GAT	CDR3 Y TAT
1 1	V	W TGG	09	:	Ħ	85 T ACG	H
 	R AGG	A GCC	9	:	н	999	CAA
	D GAC 2	 V GTA (	IMGT .	:	ı	S TCT (	105 Q CAG (
i !	G GGA	.	<b> </b>	:	m	:	C C TIGIT
i !	15 V GTA (	, i	CDR2	•	ជ	•	Y TAC 7
E+	S TCA C		İ	S	ſτι	80 G GGA	Y
ტ	T ACT	:	55	A GCA		S AGT (	V GTT
Σ	S	H. :		S TCG (		ဗ ဗဗ္ဗ	100 A GCA (
н	M ATG	- IMGT 0	Ŷ	Y TAC	 	T ACT 0	L CTG
1	10 F TTC 2	CDR1 - 30 A	i !	I ATT	1	F TTC 7	D GAC (
Н	K AAA	T CI		L CTG	1	75 R CGC 7	E GAA (
pc;	H CAC	S AGT	T 20	L CTA (		D GAT (	A GCT (
Ēι	S TCT (	V GTG 2	r. O	K AAA (	; ; ;	:	95 Q CAG (
!	O CAG	AT		P CCT 7	1	F S	V GTG 0
	5 T ACG (	25 Q CAG G	Ħ	S TCT (		V GTC C	S AGT G
	M ATG A	S S AGT C		O CAA 1	1	70 G GGA G	S AGC A
1	V GTA A	A GCC A	45	GGA C	1	T ACT G	I ATC A
 	I ATC G	K AAG G		P CCA G	 	Y TAC A	90 T ACC A
>	1 D GAC A	C TGC A	ቹ <i>እ</i>	K AAA G	1	R CGG T	F. TTC A

P L T F G A G CCG CTC ACG TTC GGT GCT GGG



FIG. 63A

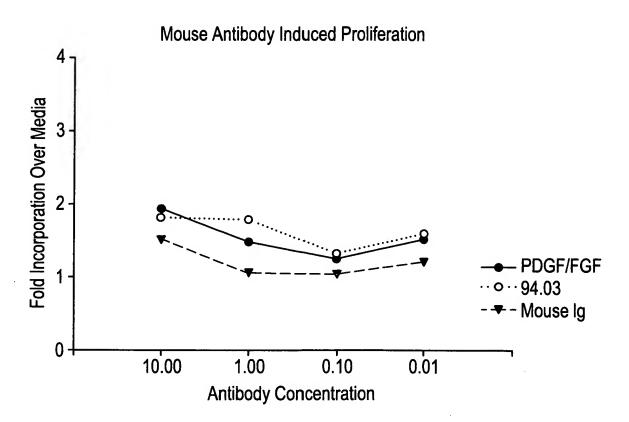


FIG. 63B

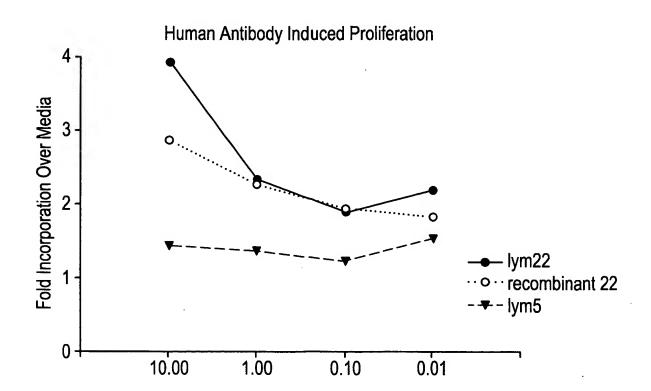
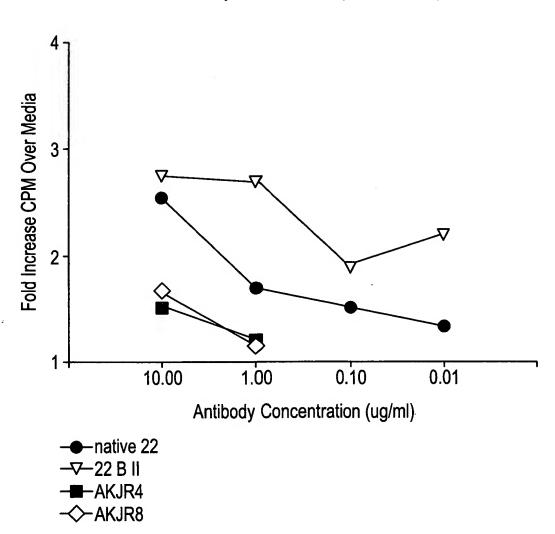




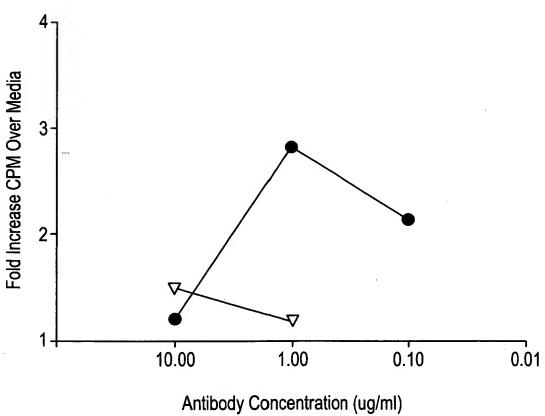
FIG. 64

### Human Antibody Induced 3H Thymidine Incorporation

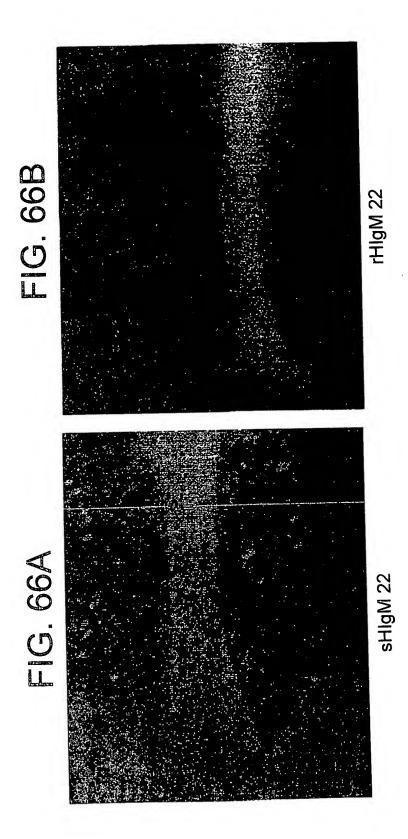




Mouse Antibody Induced 3H Thymidine Incorportion



● 09 ▼ SHL Ig



PYTFGGGGGGGCGGAGAATAAACGG



# TRANSLATION OF OI KAPPA CHAIN

FIG. 68

	N AAT	E GAG	-	TACT		TACT	IMGT 110 ' Y 'A TAC
İ	I ATT	CAA	v	65	H	FTC	- IN E GAA
	20 T ACT	Y		:	ტ	D GAT	CDR3 N AAT
	I ATT	¥ TGG		:	Σ	85 T ACA	H CAT
	T ACC	40 A GCC		:	Н	G GGT	CAG
	E GAA	L L TTA	E	:	1	S TCT (	105 O CAA
	GGA (		- IMGT	09	m	•	V DET
1	15 P CCT		CDR2		<b>K</b>		Y TAC
E	STCT		Ü	S	ſщ	80 GGA	Y
ტ	A GCA	35		g GGA		S AGT (	A M
Σ	A GCT (	IMGT		S		5 9	100 A GCA
н	L	н !	Ŷ	55 Y TAC		S AGT (	FTTT
ı	10 Y TAT	CDR1 Y TAT		I ATC		TIC	D
<del>~</del>	s TCT '	K AAA		CTT		75 R AGG	E GAA (
· 🕰	P CCA '	30 S AGC	i E	CIT	1	S	FOO HOO
, <b>E</b> 4	S TCT	IATT	ა ტ	K AAG	1	•	95 E GAG
-	O CAG	S	Σ	50 N AAT		CCA	L
1	5 T ACC (	K	· H	TACT		I ATT	S AGC (
į	I ATA	S SAGT	ı	K AAA	!	70 G GGA 7	S AGT 7
	Q CAG	25 A GCA /		999		s TCT (	I ATC /
	V GTC (	R AGG (	_α	P CCT (	İ	Q CAA 1	90 T ACC A
;	1 D GAT (	C C TIEC N	[tri	45 K AAA (	 	r TTG (	L L CIC #
•	0	· .		7 124	•	C C	



# TRANSLATION OF HNK-1 KAPPA CHAIN

	TACT	Q CAG	S S T L	S I CT	IMGT 110
	r CTC	O	· 65	T Y TAT	Σ <u>i</u> . Η (
	20 S AGT	L		GAT	CDR3
	V GTC	TGG		M 85 S TCA	
	r Aga	40 N AAC		. I 599	
	e gaa	L TTA	IMGT	s l	105
	G GGA		. O9	. m	<b>`</b>
1	15 L CTG	:	CDR2	<u>~</u>	
H	S TCT	- 40	0 8 E	F F 80 R AGG	
ტ	A GCC	35	E A	s SAGT	
Σ	s TCT	IMGT	<b>₹</b>	) ပြုပြု	100
н	L TTA	1 :	. V. 55	S	
ı	10 S TCC	CDR1 S AGC	HA	TF TTC	
٦	S	S AGT	i F	75 75 AGG	
ĸ	P	30 G	ا « در	AAA AAA	
ſτι	S TCT	IATT	0 A	:	95
1	Q CAG	D	Μ 50 1 1	200 400	
	5 T ACC	Q CAG	_ F	, ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' ' '	
	M ATG	25 A S GCA AGT	ט פ	70 0 0 0 0	į
	Q CAG	25 A GCA	2 D D	ST	
	I Q ATC CAG	R CGG	מינ	D S D S GAT TCT	06
>	1 D GAC	CTGT	F R 2 45 E P D	TI ITA	

L Q Y A S CTA CAA TAT GCT AGT ၁ TGT Y Y TAT D GAC V GTA F TTT D GAT E GAA s TCT E GAG CTT L T I S S CTC ACC ATC AGC AGC

P Y T F G G G T K L E I K R CCG TAC ACG TTC GGA GGG GGG ACC AAG CTG GAA ATA AAA CGG



## TRANSLATION OF A2B5 KAPPA CHAIN

1	S TCC	Q CAG	z	D4:	s TCT	IMGT 110 S Y GT TAC	:
	I ATA T	 Q Q CAG C	<b>V</b>	ad i : ₽	Y TAC T	- IMG' S AGT T	
E		1	65	: :	S ; TCT T2	ı	
ტ	20 T ACC	Y TAC		: 0		U.	
	V GTC	W TGG		Σ	85 T ACC	Y	
Σ	K AAG	40 Y TAC		: н	999	O CAG	
Н	E GAG	M ATG	IMGT -	: 1	STCT	105 Q CAG	
1	9 9		1 09	: m	: '	ု ပည္	
	15 P CCA		CDR2	: ~	:	Y TAC	
ፚ	s TCT	:	w	TCC F	80 6 666	Y TAT	R GGG
  -  -	A GCA	35	E	ACA .	s AGT	TACT	K AAA
	STCT	IMGT	rc.	သဗ္ဗာ	ဗ ဗ	100 A GCC	L CTG
1	M ATG	i ·	√. \.	TAT	S AGT	A GCT	e Gag
1	IATC	CDR1	<b>H</b>	ATT	F TTC	D	120 L CTG
1	10 A GCA	Y TAC	×	TGG	75 R CGC	E GAA	K AAG
1	P	30 s AGT	' ⊡	222	A GCT	A GCT	T ACC
	s TCT	V GTA	o S	AAA	:	95 E GAG	ტ ტტტ
	Q CAG	S AGT	M 250	ည	P CCT	M ATG	A GCT
	5 T ACC	S	ы	100	V GTC	s AGC	115 G GGT
1	LCTC	S AGC	, ω	100	70 G GGA	S AGC	F
1	V GTT	25 A GCC	2	GGA	S	IATC	TACG
!!!	I ATT	S	<del>с.</del>	CCA	A GCT	90 T ACA	CHC
<b>+</b>	1 O CAA	L L	면 4 건 전	AAG	L CTG	L CIO	P



LYM 46 Heavy Chain Sequence:

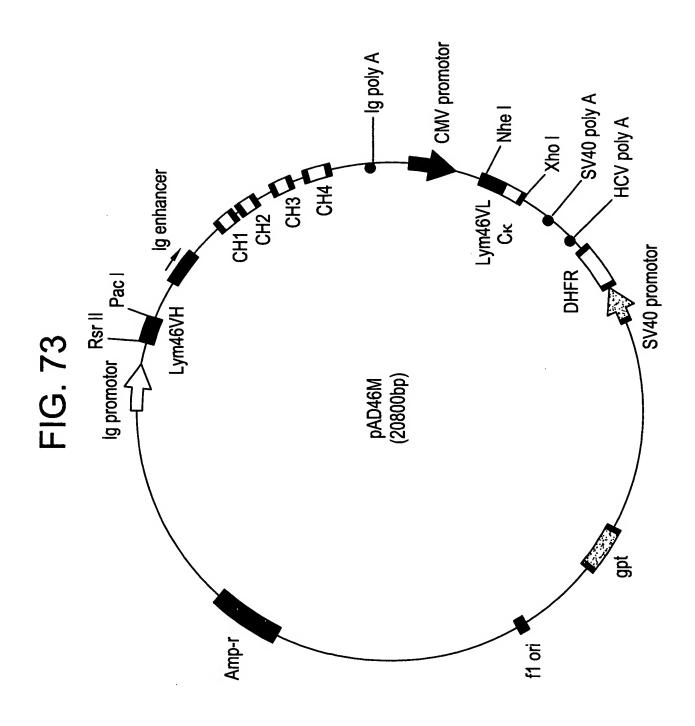
			FI	R 1										
E	V	Q	L	v	E	S	G	G	G	L	V	Q	P	G
GAG	GTG	CAG	CTG	GTG	GAG	TCT	GGG	GGA	GGC	TTG	GTC	CAG	CCT	GGG
												CDI	R1	
							A							
GGG	TCC	CTG	AGA	CTC	TCC	TGT	GCA	GCC	TCT	GGA	TTC	ACC	TTT	AGT
							ਜਾਸ 2	,						
<u>s</u>	Y	W	M	T	W	v	_FR 2 R		A	P	G	ĸ	G	
							CGC							
									,				-	
		-						CI	DR2			<del></del>		
							K							
GAG	TGG	GTG	GCC	AAC	ATA	AAG	AAA	GAT	GGA	AGT	GAG	AAA	TCC	TAT
		?R3												
							F							
GTG	GAC	TCT	GTG	AAG	GGC	CGA	TTC	ACC	ACC	TCC	AGA	GAC	AAC	GCC
K	N	s	L	Y	L	0	M	N	S	L	R	A	E	D
							ATG							
						0.0			CDI	R3			_	
							R							
ACG	GCT	GTG	TAT	TAC	TGT	GCG	AGA	CCC	AAT	TGT	GGT	GGT	GAC	TGC
- <del>-</del>	Т.	P	w	V	F	<u></u>	L	w	G	D	G	TT.	<b>T</b> .	37
TAT	TTA	CCA	TGG	ተ ጥ <u>ል</u> ሮ	ጥጥር	CAT	CTC	ጥ <u>ር</u> ር	GGC	ССШ ₹/	GGC	y C C	CILC:	C⊞C V
	~	~~~	700	420		JAL		193	360	CGT	335	ACC	CIG	GIC
T	v	s	s											

P Q A F G Q G T K V E I K R T V A P S V F CT CAG GCG TTC GAG GTG GAA ATC AAA CGA ACT GTG GCT GCA TCT GTC TTC

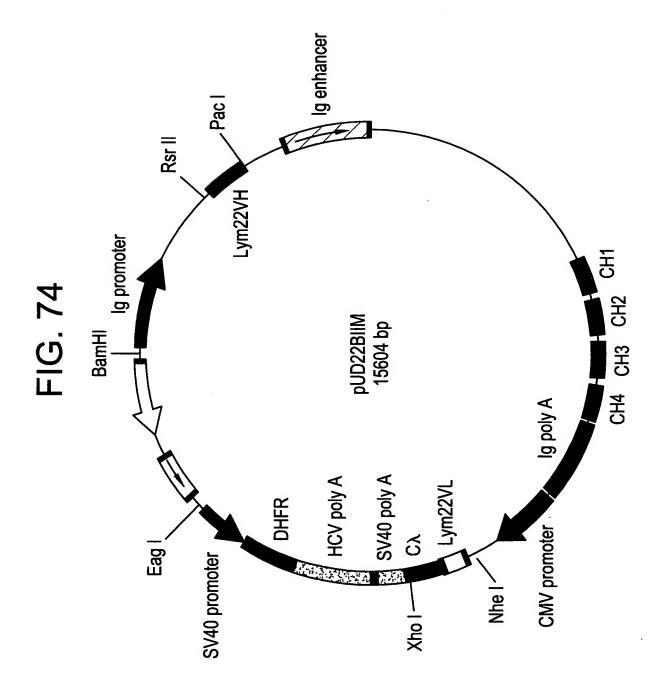


ATENT B							
	1	N AAC	CAG	<b>A</b> CC		ACT	IMGT 110 T T
	!	I ATC	CAG	· 63 :	EH	F	- IIV NAT
	* ! !	20 FCC	Y TAC	:	ტ	D GAT	CDR3 Y TAT
		<b>₹</b>	T G G	:	Σ	85 7 ACA	HAT TAT
		R AGG	40 A GCT	:	н	ა ე	O'S
		e Gag	TTP	IMGT	1	STCT	105 CAG
	!	စ ပစ္	TAC		ო		v Dar
		15 L CTG	AAC	CDR2	<b>14</b>	•	¥ TAC
	H	STCT	A A A G	Igw	Œ	80 0 0 0	YAT TAT
72	ტ	V GTG	N NAT	<b>₹</b> 36		S AGC	OFF.
<u>ල</u>	Σ	GCT	IMGT	# £ £		<b>ဗ</b> ဗ္ဗ	100 420 603
<u> </u>	н	CIG	I N II	55 TAC		S AGT	9 tg
SEQUENCE	ı	10 10 100	CDR1	ATT	!	TTC	G A D
nõas	н	DGAC	TAC	CTC		75 R CGA	ы <b>б</b>
	ĸ	4 5 5 7	30 L TTA	GE C		GAC	¢ SCT
CHAIN	Ēų	STCT	ort	G AAA		•	9 0 0 0 0 0 0
	:	CAC	SAGT	M 50 P CCT		PCCT	។ ឯ
46 KAPPA LIGHT		5 ACC	CAG	I P CCT	į	orc Grc	က နှ
PPA		M ATG	AGC	o CAG		70 G G G G	8 & C
KAE		V GTG	25 S TCC	2 6 6 8	į	a TCC	ATC
46		H ATC	K AAG	ж С Р	!	GAA	06 K
MX'		D Sec	ပ ပည်	· wkk	-	ત્ર <u>છે</u>	크 된 한

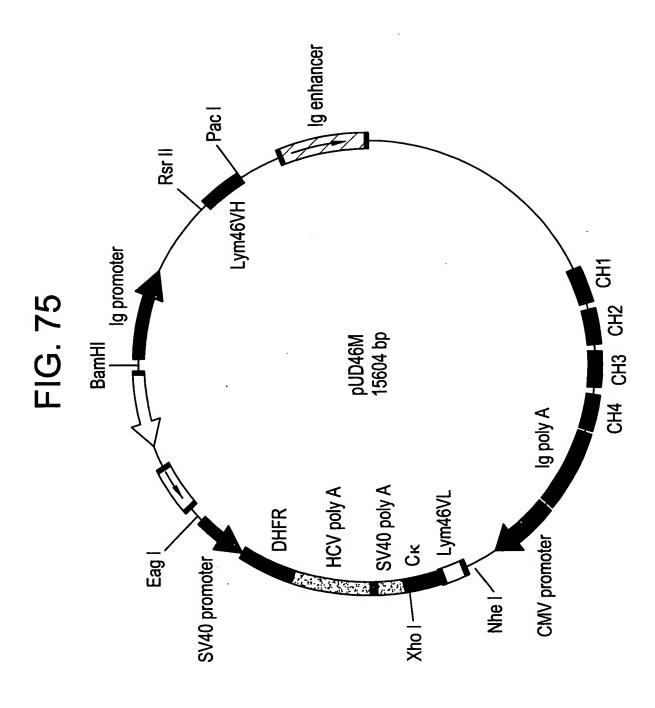




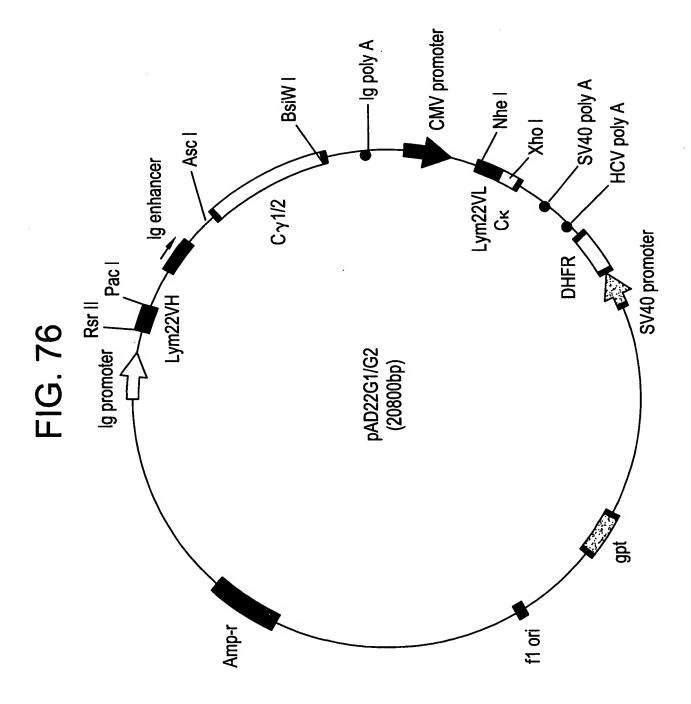




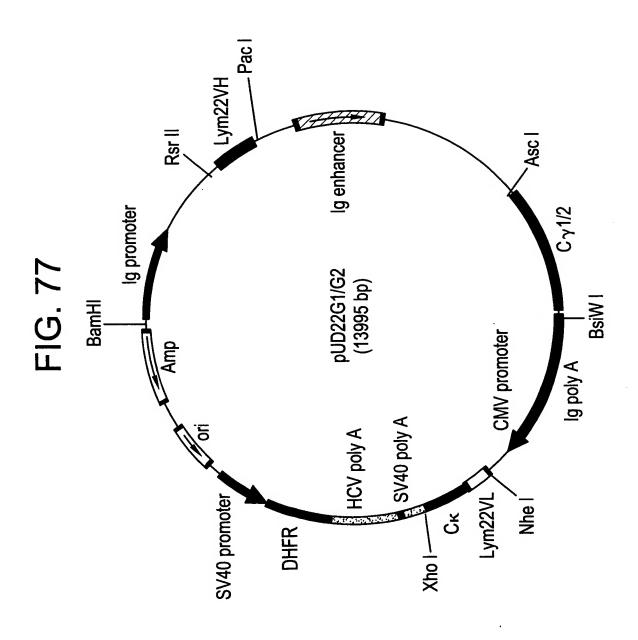




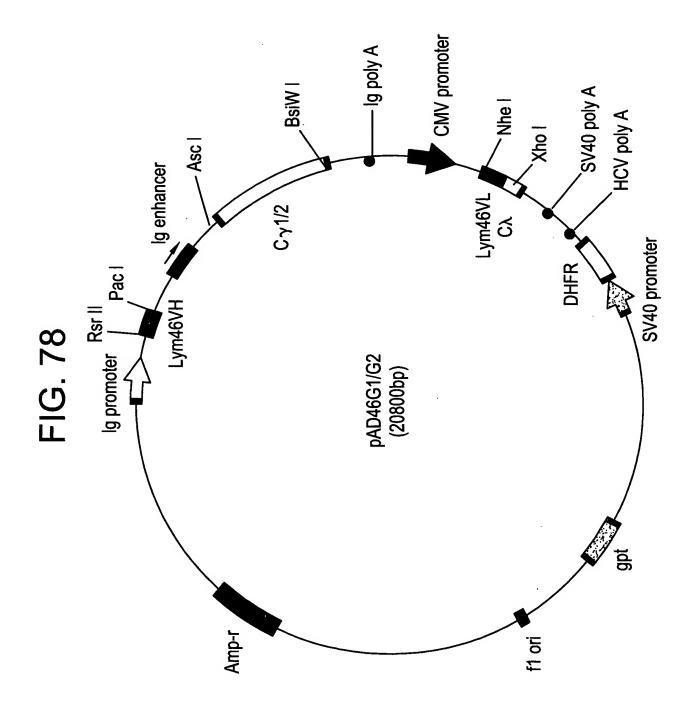




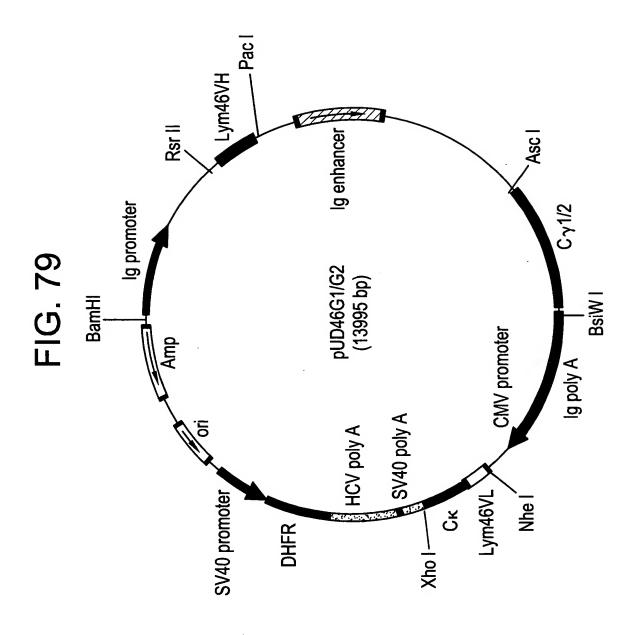










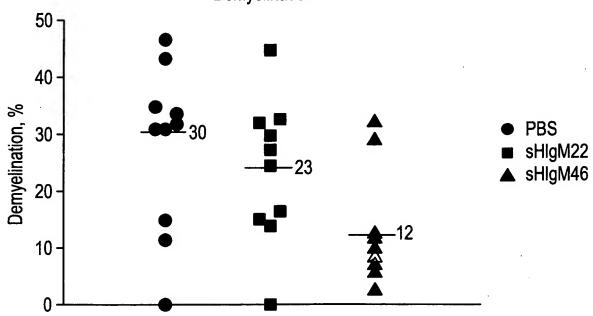




### FIG. 80A

TMEV Infected SJL Mice
Treated at 21 Days Post Infection

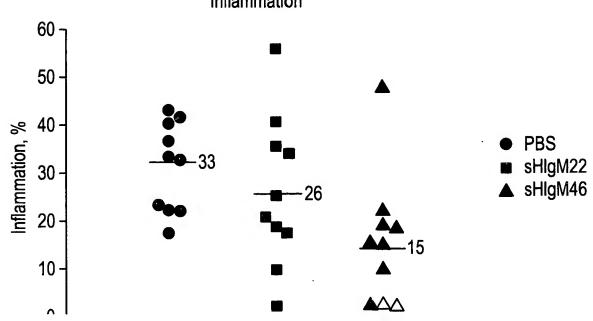
Demyelination



### FIG. 80B

TMEV Infected SJL Mice
Treated at 21 Days Post Infection

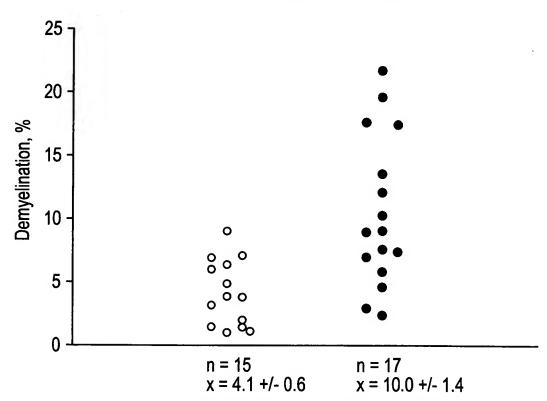
Inflammation







Chronically TMEV Infected SJL Mice Treated with sHIgM46 or sHIgM22



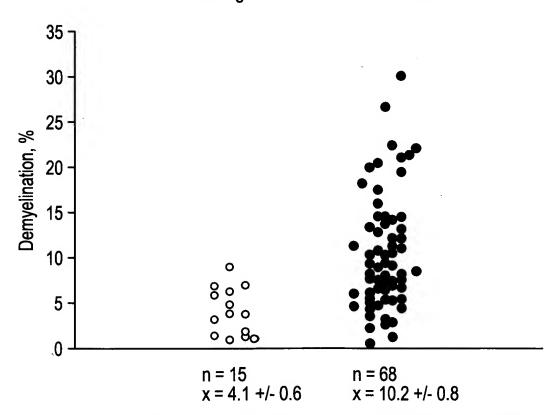
Groups are different by ANOVA, P = 0.001

- o sHlgM46
- sHlgM22



FIG. 82

Chronically TMEV Infected SJL Mice Treated sHIgM46 vs All Other Antibodies

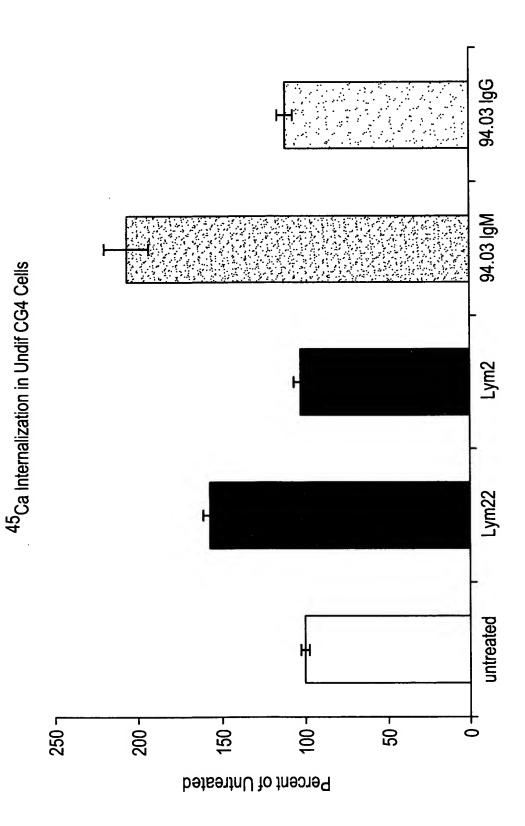


Groups are different by one way ANOVA, P = < 0.001

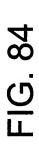
- o sHlgM46
- other mAbs

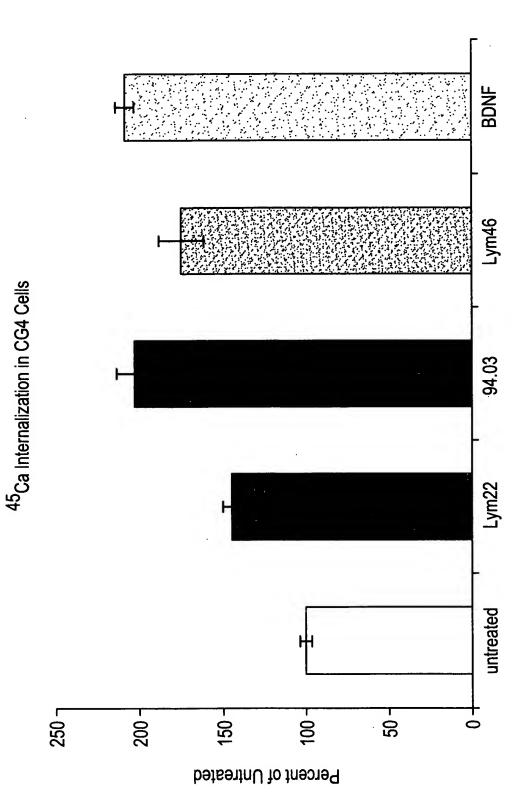


FIG. 83



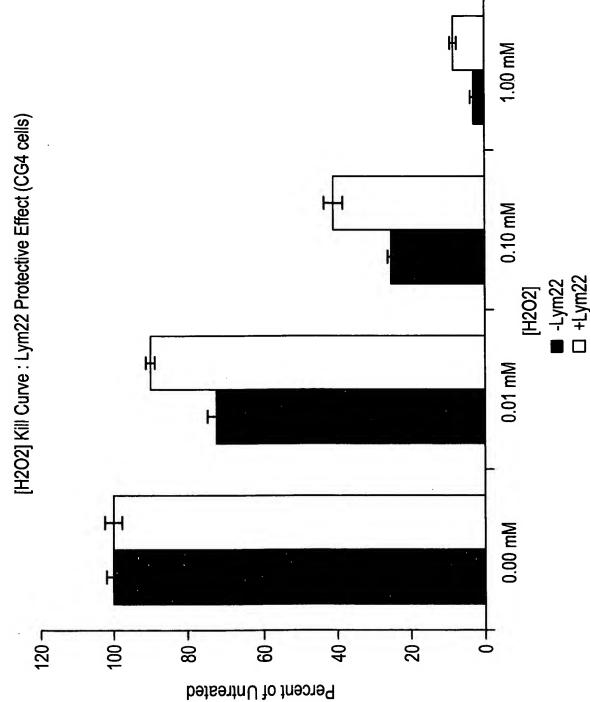






O ST 25 TOTE &

FIG. 85





### FIG. 86A

MTT Assay: H₂O₂-induced cell death

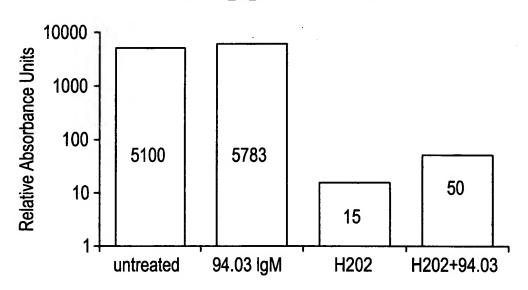


FIG. 86B

Cell Number: H₂O₂-induced cell death

